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## INTERNATIONAL APPLICATION PUBLISHED UNDER THE PATENT COOPERATION TREATY (PCT)

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<b>(21) International Application Number:</b> PCT/EP99/04698 <b>(22) International Filing Date:</b> 28 June 1999 (28.06.99)  <b>(30) Priority Data:</b> 98/08777                      6 July 1998 (06.07.98)                      FR  <b>(71) Applicants (for all designated States except US):</b> MERIAL [FR/FR]; 17, rue Bourgelat, F-69002 Lyon (FR). THE QUEEN'S UNIVERSITY OF BELFAST [GB/GB]; Stoney Road, Stormont, Belfast BT4 3SD (GB). UNIVERSITY OF SASKATCHEWAN [CA/CA]; 52 Campus Drive, Saskatoon, Saskatchewan S7W 5B4 (CA).  <b>(72) Inventors; and</b> <b>(75) Inventors/Applicants (for US only):</b> ALLAN, Gordon, Moore [GB/GB]; 51 Cabinhill Gds, Belfast BT5 7AR (GB). MEEHAN, Brian, Martin [GB/GB]; 26 St John's Close, 2 Laganbank Road, Belfast BT1 3LX (GB). ELLIS, John, Albert [US/CA]; 812, 13th Street East, Saskatoon, Saskatchewan S7N 0M3 (CA). KRAKOWKA, George, Steven [US/US]; 2676 Summit Street, Columbus, OH 43202 (US). AUDONNET, Jean-Christophe, Francis [FR/FR]; 119, rue de Créqui, F-69006 Lyon (FR).		<b>(74) Agent:</b> MONCHENY, Michel; Cabinet Lavoix, 9, place d'Estienne d'Orves, F-75441 Paris Cedex 09 (FR).  <b>(81) Designated States:</b> AE, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, CA, CH, CN, CU, CZ, DE, DK, EE, ES, FI, GB, GD, GE, GH, GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MD, MG, MK, MN, MW, MX, NO, NZ, PL, PT, RO, RU, SD, SE, SG, SI, SK, SL, TJ, TM, TR, TT, UA, UG, US, UZ, VN, YU, ZA, ZW, ARIPO patent (GH, GM, KE, LS, MW, SD, SL, SZ, UG, ZW), Eurasian patent (AM, AZ, BY, KG, KZ, MD, RU, TJ, TM), European patent (AT, BE, CH, CY, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE), OAPI patent (BF, BJ, CF, CG, CI, CM, GA, GN, GW, ML, MR, NE, SN, TD, TG).  <b>Published</b> <i>Without international search report and to be republished upon receipt of that report.</i>
<b>(54) Title:</b> PORCINE CIRCOVIRUS AND PARVOVIRUS VACCINE		
<b>(57) Abstract</b> <p>The invention relates to antigenic preparations and vaccines directed against the porcine multisystemic wasting syndrome (PMWS), comprising at least one porcine circovirus antigen, preferably type II, and at least one porcine parvovirus antigen.</p>		

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"Porcine circovirus and parvovirus vaccine"

The present invention relates to a vaccine against the PMWS syndrome (*Porcine Multisystemic Wasting Syndrome* also called *Post-Weaning Multisystemic Wasting Syndrome*).

5 Various documents are cited in the following text, and various documents are referenced or cited in documents cited in the following text. There is no admission that any of these documents are indeed prior art as to the present invention. All documents cited  
10 herein and all documents referenced or cited in documents cited herein are hereby incorporated herein by reference.

PCV (for "Porcine CircoVirus") was originally detected as a noncytopathogenic contaminant in pig  
15 kidney cell lines PK/15. This virus was classified among the Circoviridae with the chicken anaemia virus (CAV for *Chicken Anaemia Virus*) and the PBFDV virus (*Pscittacine Beak and Feather Disease Virus*). It is a small nonenveloped virus (from 15 to 24 nm) whose  
20 common characteristic is to contain a genome in the form of a circular single-stranded DNA of 1.76 to 2.31 kb. It was first thought that this genome encoded a polypeptide of about 30 kDa (Todd et al., Arch Virol 1991, 117; 129-135). Recent work has however shown a  
25 more complex transcription (Meehan B. M. et al., 1997, 78; 221-227). Moreover, no significant homologies in nucleotide sequence or in common antigenic determinants are known between the three types of circoviruses known.

30 The PCV derived from the PK/15 cells is considered not to be pathogenic. Its sequence is known from B.M. Meehan et al., J. Gen. Virol 1997 (78) 221-227. It is only very recently that some authors have thought that strains of PCV could be pathogenic  
35 and associated with the PMWS syndrome (Gupi P.S. Nayar et al., Can. Vet. J, vol. 38, 1997: 385-387 and Clark E.G., Proc. Am. Assoc. Swine Prac. 1997; 499-501). Nayar et al. have detected PCV DNA in pigs having the PMWS syndrome using PCR techniques.

The PMWS syndrome detected in Canada, the United States and France is clinically characterized by a gradual loss of weight and by manifestations such as tachypnea, dyspnea and jaundice. From the pathological point of view, it is manifested by lymphocytic or granulomatous infiltrations, lymphadenopathies and, more rarely, by hepatitis and lymphocytic or granulomatous nephritis (Clark E.G., Proc. Am. Assoc. Swine Prac. 1997; 499-501; La Semaine Vétérinaire No. 26, supplement to La Semaine Vétérinaire 1996 (834); La Semaine Vétérinaire 1997 (857): 54; Gupi P.S. Nayar et al., Can. Vet. J, vol. 38, 1997; 385-387).

The applicant has succeeded in isolating five new PCV strains from pulmonary or ganglionic samples obtained from farms situated in Canada, the United States (California) and France (Brittany). These viruses have been detected in lesions in pigs with the PMWS syndrome, but not in healthy pigs.

The applicant has, in addition, sequenced the genome of four of these strains, namely the strains obtained from Canada and the United States as well as two French strains. The strains exhibit a very strong homology with each other at the nucleotide level, exceeding 96% and much weaker with the PK/15 strain, about 76%. The new strains can thus be considered as being representative of a new type of porcine circovirus, called here type II, type I being represented by PK/15.

Purified preparations of five strains were deposited under the Budapest Treaty at the ECACC (European Collection of Cell Cultures, Centre for Applied Microbiology & Research, Porton Down, Salisbury, Wiltshire SP4 0JG, United Kingdom) on Thursday 2 October 1997:

- accession No. V97100219 (called here Imp. 1008PCV)
  - accession No. V9700218 (called here Imp. 1010PCV)
  - accession No. V97100217 (called here Imp. 999PCV),
- and, on Friday 16 January 1998:
- accession No. V98011608 (called here Imp. 1011-48285)

- accession No. V98011609 (called here Imp. 1011-48121).

5 The applicant has observed that, in a trial for experimental reproduction of the porcine multisystemic wasting syndrome, a porcine parvovirus combined with the porcine circovirus could lead to a worsening of the disease.

10 The subject of the present invention is therefore a vaccination of pigs using a porcine circovirus, in particular type I or type II, preferably type II, vaccine, combined with a vaccination with a porcine parvovirus vaccine. This is understood to mean vaccination with either a bivalent vaccine, or the simultaneous use, in pigs, of a porcine circovirus vaccine and of a porcine parvovirus vaccine.

15 The reference parvovirus strain is the NADL-2 strain which is accessible from the ATCC collection under the reference VR-742. Vaccination against the porcine parvovirus is well known to persons skilled in the art and vaccines against the porcine parvovirus are commercially available. There may be mentioned by way of example: Parvovax® (inactivated vaccine against porcine parvoviro-sis, distributed by MERIAL). See also e.g. P. Vannier et A. Laval., Point. Vét. 1993, 25  
20 (151), 53-60 ; G. Florent et al., Proceedings of the Ninth Congress of Pig Veterinary Society, July 15-18, 1986, Barcelona, Spain. For DNA vaccines, one can refer e.g. to WO-A-98 03658.

30 The subject of the present invention is therefore an antigenic preparation directed against the PMWS syndrome, comprising at least one porcine circovirus antigen (preferably type II circovirus) and at least one porcine parvovirus antigen. In accordance with the invention, the porcine circovirus antigen (preferably  
35 type II circovirus) and the porcine parvovirus antigen comprise, independently of each other, an antigen chosen from the group consisting of an attenuated live whole antigen, an inactivated whole antigen, a subunit antigen, a recombinant live vector and a DNA vector. It

is understood that the combination according to the invention may involve the use of any appropriate antigen or antigenic preparation form, it being understood that it is not necessary to use the same form for a given combination. The antigenic preparation may comprise, in addition, as is known per se, a vehicle or excipient acceptable from the veterinary point of view, and optionally an adjuvant acceptable from the veterinary point of view.

10 The subject of the present invention is also an immunogenic composition or a vaccine against the PMWS syndrome, comprising an effective quantity of circovirus + parvovirus antigenic preparation as described above, in a vehicle or excipient acceptable from the veterinary point of view, and optionally an adjuvant acceptable from the veterinary point of view. An immunogenic composition elicits an immunological response which can, but need not be, protective. A vaccine composition elicits a protective response. Accordingly, the term "immunogenic composition" include a vaccine composition" (as the former term can be protective composition).

The subject of the invention is also an immunological or a vaccination kit containing, packaged separately, an antigenic preparation or an immunogenic composition or a vaccine against the porcine circovirus and an antigenic preparation or an immunogenic composition or a vaccine against the porcine parvovirus. This kit may have the various characteristics set out above for the antigenic preparations, immunogenic compositions and vaccines.

The subject of the invention is also a method of immunization or of vaccination against the PMWS syndrome, comprising the administration of an immunogenic composition or a vaccine against the porcine circovirus and of an immunogenic composition or a vaccine against the porcine parvovirus or the administration of a bivalent immunogenic composition or vaccine, comprising, in the same formulation, an



antigenic preparation specific to each virus. This method of immunisation or vaccination uses in particular the vaccines as defined above.

5 The subject of the invention is also the use of an antigenic preparation or of an immunogenic composition or a vaccine against the parvovirus, as in particular defined supra, for the preparation of a pharmaceutical composition intended to be used in the context of the prevention of the PMWS syndrome, in  
10 combination with an antigenic preparation or an immunogenic composition or a vaccine against the porcine circovirus.

For the production of circovirus antigenic preparations, the circoviruses may be obtained after  
15 passage on cells, in particular cell lines, e.g. PK/15 cells. The culture supernatants or extracts, optionally purified by standard techniques, may be used as antigenic preparation.

In the context of attenuated antigenic preparations and attenuated immunogenic compositions or vaccines, the attenuation may be carried out according to the customary methods, e.g. by passage on cells, preferably by passage on pig cells, especially cell  
20 lines, such as PK/15 cells (for example from 50 to 150, especially of the order of 100, passages). These immunogenic compositions and vaccines comprise in general a vehicle or diluent acceptable from the veterinary point of view, optionally an adjuvant acceptable from the veterinary point of view, as well  
25 as optionally a freeze-drying stabilizer.

These antigenic preparations, immunogenic compositions and vaccines will preferably comprise from  $10^3$  to  $10^7$  TCID<sub>50</sub> of the attenuated virus in question.

They may be antigenic preparations, immunogenic  
35 compositions and vaccines based on inactivated whole antigen. The inactivated immunogenic compositions and vaccines comprise, in addition, a vehicle or a diluent acceptable from the veterinary point of view, with

optionally in addition an adjuvant acceptable from the veterinary point of view.

The circoviruses according to the invention, with the fractions which may be present, are inactivated according to techniques known to persons skilled in the art. The inactivation will be preferably carried out by the chemical route, e.g. by exposing the antigen to a chemical agent such as formaldehyde (formalin), paraformaldehyde,  $\beta$ -propiolactone or ethyleneimine or its derivatives. The preferred method of inactivation will be herein the exposure to a chemical agent and in particular to ethyleneimine or to  $\beta$ -propiolactone.

Preferably, the inactivated antigenic preparations and the inactivated immunogenic compositions and vaccines according to the invention will be supplemented with adjuvant, advantageously by being provided in the form of emulsions, for example water-in-oil or oil-in-water, according to techniques well known to persons skilled in the art. It will be possible for the adjuvant character to also come from the incorporation of a customary adjuvant compound into the active ingredient.

Among the adjuvants which may be used, there may be mentioned by way of example aluminium hydroxide, the saponines (e.g. Quillaja saponin or Quil A; see Vaccine Design, The Subunit and Adjuvant Approach, 1995, edited by Michael F. Powel and Mark J. Newman, Plenum Press, New-York and London, p.210), Avridine® (Vaccine Design p. 148), DDA (Dimethyldioctadecylammonium bromide, Vaccine Design p. 157), Polyphosphazene (Vaccine Design p. 204), or alternatively oil-in-water emulsions based on mineral oil, squalene (e.g. SPT emulsion, Vaccine Design p. 147), squalene (e.g. MF59, Vaccine Design p. 183), or water-in-oil emulsions based on metabolizable oil (preferably according to WO-A-94 20071) as well as the emulsions described in US-A-5,422,109. It is also

possible to choose combinations of adjuvants, for example Avridine® or DDA combined with an emulsion.

These antigenic preparations, immunogenic compositions and vaccines will preferably comprise from  
5  $10^5$  to  $10^8$  TCID<sub>50</sub> of the inactivated whole virus in question.

The adjuvants for live vaccines described above can be selected from those given for the inactivated. The emulsions are preferred. To those indicated for the  
10 inactivated vaccine, there may be added those described in WO-A-9416681.

As freeze-drying stabilizer, there may be mentioned by way of example SPGA (Bovarnik et al., J. Bacteriology 59, 509, 950), carbohydrates such as  
15 sorbitol, mannitol, starch, sucrose, dextran or glucose, proteins such as albumin or casein, derivatives of these compounds, or buffers such as alkali metal phosphates.

The antigenic preparations, immunogenic compositions and vaccines according to the invention may comprise one or more active ingredients (antigens) of one or more circoviruses and/or parvoviruses according to the invention.

The applicant has, in addition, obtained the  
25 genome of four of the type II porcine circovirus isolates, identified SEQ ID NO: 1 to 4. The sequence of strain PK-15 is given as SEQ ID NO: 5. It goes without saying that the invention automatically covers the equivalent sequences, that is to say the sequences  
30 which do not change the functionality or the strain-specificity of the sequence described or of the polypeptides encoded by this sequence. There will of course be included the sequences differing by degeneracy of the code.

35 The invention also covers the equivalent sequences in the sense that they are capable of hybridizing with the above sequence under high stringency conditions and/or have a high homology with the strains of the invention.

These sequences and their fragments can be advantageously used for the in vitro or in vivo expression of polypeptides with the aid of appropriate vectors.

5 In particular, the open reading frames (ORF1-13), forming DNA fragments according to the invention, which can be used to this effect have been identified on the genomic sequence of the type II circoviruses. The invention relates to any polypeptide containing at  
10 least one of these open reading frames (corresponding amino acid sequence). Preferably, the invention relates to a protein essentially consisting of ORF4, ORF7, ORF10 or ORF13.

For the expression of subunits in vitro, as a  
15 means of expression, *E. coli* or a baculovirus will be preferably used (US-A-4,745,051). The coding sequence(s) or their fragments may be integrated into the baculovirus genome (e.g. the baculovirus *Autographa californica* Nuclear Polyhedrosis Virus AcNPV) and the  
20 latter can be then propagated on insect cells, e.g. *Spodoptera frugiperda* Sf9 (deposit ATCC CRL 1711). The subunits can also be produced in eukaryotic cells such as yeasts (e.g. *Saccharomyces cerevisiae*) or mammalian cells (e.g. CHO, BHK).

25 The subject of the invention is also the use as subunits of the polypeptides which will be produced in vitro by these expression means, and then optionally purified according to conventional techniques. The subunit immunogenic compositions and vaccines comprise  
30 at least one polypeptide as thus obtained, or fragment, in a vehicle or diluent acceptable from the veterinary point of view and optionally an adjuvant acceptable from the veterinary point of view.

For the expression in vivo for the purpose of  
35 producing immunogenic compositions and vaccines of the recombinant live type or DNA type, the coding sequence(s) or their fragments are inserted into an appropriate expression vector under conditions allowing the expression of the polypeptide(s). As appropriate

live vectors, there may be used preferably live viruses, preferably capable of multiplying in pigs, nonpathogenic for pigs (naturally nonpathogenic or rendered as such), according to techniques well known to persons skilled in the art. There may be used in particular pig herpesviruses such as Aujeszky's disease virus, porcine adenovirus, poxviruses, especially vaccinia virus, avipox virus, canarypox virus, swinepox virus. DNA vectors can also be used as vectors (WO-A-9011092, WO-A-9319813, WO-A-9421797, WO-A-9520660).

The subject of the invention is therefore also the vectors and the recombinant live type or DNA (polynucleotide) type immunogenic compositions or vaccines thus prepared, their preparation and their use, the immunogenic compositions and the vaccines comprising, in addition, a vehicle or diluent acceptable from the veterinary point of view.

By definition, a DNA immunogenic composition or vaccine comprises a DNA vector which is a circular vaccinal plasmid, supercoiled or otherwise, or a linear DNA molecule, incorporating and expressing in vivo a nucleotide sequence encoding an antigenic polypeptide.

Recombinant and DNA-type immunogenic compositions and vaccines may comprise an adjuvant.

In the context of the combined immunization or vaccination programmes, it is also possible to combine the immunization or vaccination against the porcine circovirus and the porcine parvovirus with a an immunization or vaccination against other pig pathogens, in particular those which could be associated with the PMWS syndrome. The immunogenic composition or vaccine according to the invention may therefore comprise another valencey corresponding to another pig pathogen chosen from PRRS (Porcine Reproductive and Respiratory Syndrome) and/or Mycoplasma hyopneumoniae, and/or E. coli, and/or Atrophic Rhinitis, and/or Pseudorabies (Aujeszky's disease) virus and/or porcine influenza and/or Actinobacillus pleuropneumoniae and/or Hog cholera, and

combinations thereof. Preferably, the programme of immunization or vaccination and the vaccines according to the invention will combine immunizations or vaccinations against the circovirus and the parvovirus, and the PRRS (WO-A-93/07898, WO-A-94/18311, FR-A-2 709 966 ; C. Charreyre et al., Proceedings of the 15<sup>th</sup> IPVS Congress, Birmingham, England, 5-9 July 1998, p 139 ; and/or Mycoplasma hyopneumoniae (EP-A-597 852, EP-A-550 477, EP-A571 648 ; O. Martinon et al. p 157, 284, 285 and G. Reynaud et al., p 150, all in the above-referenced Proceedings of the 15<sup>th</sup> IPVS Congress) and/or porcine influenza. It is thus possible to use any appropriate form of immunogenic composition or vaccine, in particular any available commercial vaccine, so as to combine it with the immunogenic composition or vaccine against the porcine circovirus and porcine parvovirus as described here.

The subject of the present invention is therefore also multivalent immunogenic compositions and vaccines, multivaccine kits, and combined immunization or vaccination methods which make it possible to use such combined immunization or vaccination programmes.

The invention will now be described in greater detail with the aid of nonlimiting exemplary embodiments, taken with reference to the drawing, in which:

Figure 1: DNA sequence of the genome of the Imp. 1011-48121 strain

Figure 2: DNA sequence of the genome of the Imp. 1011-48285 strain

Figure 3: DNA sequence of the genome of the Imp. 999 strain

Figure 4: DNA sequence of the genome of the Imp. 1010 strain

Figure 5: Alignment of the 4 sequences according to Figures 1 to 4 with the sequence of the PCV PK/15 strain

## Sequence listing SEQ ID

- SEQ ID No: 1 DNA sequence of the genome of the  
Imp. 1011-48121 strain
- 5 SEQ ID No: 2 DNA sequence of the genome of the  
Imp. 1011-48285 strain
- SEQ ID No: 3 DNA sequence of the genome of the  
Imp. 999 strain
- SEQ ID No: 4 DNA sequence of the genome of the  
Imp. 1010 strain
- 10 SEQ ID No: 5 DNA sequence of the genome of the PK/15  
strain

## EXAMPLES

- 15 Example 1: Culture and isolation of the porcine  
circovirus strains:

Tissue samples were collected in France, Canada  
and the USA from lung and lymph nodes of piglets. These  
piglets exhibited clinical signs typical of the post-  
weaning multisystemic wasting syndrome. To facilitate  
20 the isolation of the viruses, the tissue samples were  
frozen at -70°C immediately after autopsy.

For the viral isolation, suspensions containing  
about 15% tissue sample were prepared in a minimum  
medium containing Earle's salts (EMEM, BioWhittaker UK  
25 Ltd., Wokingham, UK), penicillin (100 IU/ml) and  
streptomycin (100 µg/ml) (MEM-SA medium), by grinding  
tissues with sterile sand using a sterile mortar and  
pestle. This ground preparation was then taken up in  
MEM-SA, and then centrifuged at 3000 g for 30 minutes  
30 at +4°C in order to harvest the supernatant.

Prior to the inoculation of the cell cultures,  
a volume of 100 µl of chloroform was added to 2 ml of  
each supernatant and mixed continuously for 10 minutes  
at room temperature. This mixture was then transferred  
35 to a microcentrifuge tube, centrifuged at 3000 g for 10  
minutes, and then the supernatant was harvested. This  
supernatant was then used as inoculum for the viral  
isolation experiments.

All the viral isolation studies were carried out on PK/15 cell cultures, known to be uncontaminated with the porcine circovirus (PCV), pestiviruses, porcine adenoviruses and porcine parvoviruses (Allan G. et al Pathogenesis of porcine circovirus experimental infections of colostrum-deprived piglets and examination of pig foetal material. Vet. Microbiol. 1995, 44, 49-64).

The isolation of the porcine circoviruses was carried out according to the following technique:

Monolayers of PK/15 cells were dissociated by trypsinization (with a trypsin-versene mixture) from confluent cultures, and taken up in MEM-SA medium containing 15% foetal calf serum not contaminated by pestivirus (= MEM-G medium) in a final concentration of about 400,000 cells per ml. 10 ml aliquot fractions of this cell suspension were then mixed with 2 ml aliquot fractions of the inocula described above, and the final mixtures were aliquoted in 6 ml volumes in two Falcon flasks of 25 cm<sup>2</sup>. These cultures were then incubated at +37°C for 18 hours under an atmosphere containing 10% CO<sub>2</sub>.

After incubation, the culture medium of the semi-confluent monolayers were treated with 300 mM D-glucosamine (Cat # G48175, Sigma-Aldrich Company Limited, Poole, UK) (Tischr I. et al., Arch. Virol., 1987 96 39-57), then incubation was continued for an additional period of 48-72 hours at +37°C. Following this last incubation, one of the two Falcons of each inoculum was subjected to 3 successive freeze/thaw cycles. The PK/15 cells of the remaining Falcon were treated with a trypsin-versene solution, resuspended in 20 ml of MEM-G medium, and then inoculated into 75 cm<sup>2</sup> Falcons at a concentration of 400,000 cells/ml. The freshly inoculated flasks were then "superinfected" by addition of 5 ml of the corresponding lysate obtained after the freeze/thaw cycles.



Example 2: Preparation of the samples of cell culture for the detection of porcine circoviruses by immunofluorescence or by *in situ* hybridization

A volume of 5 ml of the "superinfected" suspension was collected and inoculated into a Petri dish 55 mm in diameter containing a sterile and fat-free glass coverslip. The cultures in the flasks and on glass coverslips were incubated at +37°C and treated with glucosamine as described in Example 1. The cultures on glass coverslips were harvested from 24 to 48 hours after the treatment with glucosamine and fixed, either with acetone for 10 minutes at room temperature, or with 10% buffered formaldehyde for 4 hours. Following this fixing, all the glass coverslips were stored at -70°C, on silica gel, before their use for the *in situ* hybridization studies and the immunocytochemical labelling studies.

Example 3: Techniques for the detection of PCV sequences by *in situ* hybridization

*In situ* hybridization was carried out on tissues collected from diseased pigs and fixed with formaldehyde and also on the preparations of cell cultures inoculated for the viral isolation (see Example 2) and fixed on glass coverslips.

Complete genomic probes corresponding to the PK/15 porcine circoviruses (PCV) and to the infectious chicken anaemia virus (CAV) were used. The plasmid pPCV1, containing the replicative form of the PCV genome, cloned in the form of a single 1.7 kilo base pair (kbp) insert (Meehan B. et al. Sequence of porcine circovirus DNA: affinities with plant circoviruses, J. Gen. Virol. 1997, 78, 221-227), was used as specific viral DNA source for PCV. An analogous plasmid, pCAA1, containing the 2.3 kbp replicative form of the avian circovirus CAV was used as negative control. The respective glycerol stocks of the two plasmids were used for the production and purification of the plasmids according to the alkaline lysis technique

(Sambrook J. et al. Molecular cloning: A Laboratory Manual. 2nd Edition, Cold Spring Harbor Laboratory, Cold Spring Harbor, New York, 1989) so that they are then used as templates for the preparation of the probes. The circovirus probes representative of the complete genomes of PCV and of CAV were produced from the purified plasmids described above (1 µg for each probe) and from hexanucleotide primers at random using a commercial nonradioactive labelling kit ("DIG DNA labelling kit", Boehringer Mannheim, Lewes, UK) according to the supplier's recommendations.

The digoxigenin-labelled probes were taken up in a volume of 50-100 µl of sterile water before being used for the *in situ* hybridization.

The diseased pig tissue samples, enclosed in paraffin and fixed with formaldehyde, as well as the preparations of infected cell cultures, fixed with formaldehyde, were prepared for the detection of the PCV nucleic acids according to the following technique:

Sections 5 µm thick were cut from tissue blocks enclosed in paraffin, rendered paraffin free, and then rehydrated in successive solutions of alcohol in decreasing concentrations. The tissue sections and the cell cultures fixed with formaldehyde were incubated for 15 minutes and 5 minutes respectively at +37°C in a 0.5% proteinase K solution in 0.05 M Tris-HCl buffer containing 5 mM EDTA (pH 7.6). The slides were then placed in a 1% glycine solution in autoclaved distilled water, for 30 seconds, washed twice with 0.01 M PBS buffer (phosphate buffered saline) (pH 7.2), and finally washed for 5 minutes in sterile distilled water. They were finally dried in the open air and placed in contact with the probes.

Each tissue/probe preparation was covered with a clean and fat-free glass coverslip, and then placed in an oven at +90°C for 10 minutes, and then placed in contact with an ice block for 1 minute, and finally incubated for 18 hours at +37°C. The preparations were then briefly immersed in a 2X sodium citrate salt (SSC)

buffer (pH 7.0) in order to remove the protective glass coverslips, and then washed twice for 5 minutes in 2X SSC buffer and finally washed twice for 5 minutes in PBS buffer.

5           After these washes, the preparations were immersed in a solution of 0.1 M maleic acid, 0.15 M NaCl (pH 7.5) (maleic buffer) for 10 minutes, and then incubated in a 1% solution of blocking reagent (Cat # 1096176, Boehringer Mannheim UK, Lewis, East Sussex, 10 UK) in maleic buffer for 20 minutes at +37°C.

          The preparations were then incubated with a 1/250 solution of an anti-digoxigenin monoclonal antibody (Boehringer Mannheim), diluted in blocking buffer, for 1 hour at +37°C, washed in PBS and finally 15 incubated with a biotinylated anti-mouse immunoglobulin antibody for 30 minutes at +37°C. The preparations were washed in PBS and the endogenous peroxidase activity was blocked by treatment with a 0.5% hydrogen peroxide solution in PBS for 20 minutes at room temperature. The 20 preparations were again washed in PBS and treated with a 3-amino-9-diethylcarbazole (AEC) substrate (Cambridge Bioscience, Cambridge, UK) prepared immediately before use.

          After a final wash with tap water, the 25 preparations were counterstained with hematoxylin, "blued" under tap water, and mounted on microscope glass coverslips with a mounting fluid (GVA Mount, Cambridge Bioscience, Cambridge, UK). The experimental controls included the use of a nonpertinent negative 30 probe (CAV) and of a positive probe (PCV) on samples obtained from diseased pigs and from nondiseased pigs.

#### Example 4: Technique for the detection of PCV by immunofluorescence

35           The initial screening of all the cell culture preparations fixed with acetone was carried out by an indirect immunofluorescence technique (IIF) using a 1/100 dilution of a pool of adult pig sera. This pool of sera comprises sera from 25 adult sows from Northern

Ireland and is known to contain antibodies against a wide variety of porcine viruses, including PCV: porcine parvovirus, porcine adenovirus, and PRRS virus. The IIF technique was carried out by bringing the serum (diluted in PBS) into contact with the cell cultures for one hour at +37°C, followed by two washes in PBS. The cell cultures were then stained with a 1/80 dilution in PBS of a rabbit anti-pig immunoglobulin antibody conjugated with fluorescein isothiocyanate for one hour, and then washed with PBS and mounted in glycerol buffer prior to the microscopic observation under ultraviolet light.

**Example 5: Results of the *in situ* hybridization on diseased pig tissues**

The *in situ* hybridization, using a PCV genomic probe, prepared from tissues collected from French, Canadian and Californian piglets having multisystemic wasting lesions and fixed with formaldehyde, showed the presence of PCV nucleic acids associated with the lesions, in several of the lesions studied. No signal was observed when the PCV genomic probe was used on tissues collected from nondiseased pigs or when the CAV probe was used on the diseased pig tissues. The presence of PCV nucleic acid was identified in the cytoplasm and the nucleus of numerous mononuclear cells infiltrating the lesions in the lungs of the Californian piglets. The presence of PCV nucleic acid was also demonstrated in the pneumocytes, the bronchial and bronchiolar epithelial cells, and in the endothelial cells of the arterioles, the veinlets and lymphatic vessels.

In diseased French pigs, the presence of PCV nucleic acid was detected in the cytoplasm of numerous follicular lymphocytes and in the intrasinusoidal mononuclear cells of the lymph nodes. The PCV nucleic acid was also detected in occasional syncytia. Depending on these detection results, samples of Californian pig lungs, French pig mesenteric lymph

nodes, and Canadian pig organs were selected for the purpose of isolating new porcine circovirus strains.

5      **Example 6: Results of the cell culture of the new  
porcine circovirus strains and detection by  
immunofluorescence**

10      No cytopathic effect (CPE) was observed in the  
cell cultures inoculated with the samples collected  
from French piglets (Imp.1008 strain), Californian  
15      piglets (Imp.999 strain) and Canadian piglets (Imp.1010  
strain) showing clinical signs of multisystemic wasting  
syndrome. However, immunolabelling of the preparations  
obtained from the inoculated cell cultures, after  
fixing using acetone and with a pool of pig polyclonal  
20      sera, revealed nuclear fluorescence in numerous cells  
in the cultures inoculated using the lungs of  
Californian piglets (Imp.999 strain), using the  
mediastinal lymph nodes of French piglets (Imp.1008  
strain), and using organs of Canadian piglets (Imp.1010  
strain).

**Example 7: Extraction of the genomic DNA of the porcine  
circoviruses**

25      The replicative forms of the new strains of  
porcine circoviruses (PCV) were prepared using infected  
PK/15 cell cultures (see Example 1) (10 Falcons of  
75 cm<sup>2</sup>) harvested after 72-76 hours of incubation and  
treated with glucosamine, as described for the cloning  
of the replicative form of CAV (Todd. D. et al. Dot  
30      blot hybridization assay for chicken anaemia agent  
using a cloned DNA probe. J. Clin. Microbiol. 1991, 29,  
933-939). The double-stranded DNA of these replicative  
forms was extracted according to a modification of the  
Hirt technique (Hirt B. Selective extraction of polyoma  
35      virus DNA from infected cell cultures, J. Mol. Biol.  
1967, 36, 365-369), as described by Molitor (Molitor  
T.W. et al. Porcine parvovirus DNA: characterization of  
the genomic and replicative form DNA of two virus  
isolates, Virology, 1984, 137, 241-254).

**Example 8: Restriction map of the replicative form of the genome of the porcine circovirus Imp.999 strain.**

The DNA (1-5 µg) extracted according to the Hirt technique was treated with S1 nuclease (Amersham) according to the supplier's recommendations, and then this DNA was digested with various restriction enzymes (Boehringer Mannheim, Lewis, East Sussex, UK) and the products of digestion were separated by electrophoresis on a 1.5% agarose gel in the presence of ethidium bromide as described by Todd et al. (Purification and biochemical characterization of chicken anemia agent. J. Gen. Virol. 1990, 71, 819-823). The DNA extracted from the cultures of the Imp.999 strain possess a unique EcoRI site, 2 SacI sites and do not possess any PstI site. This restriction profile is therefore different from the restriction profile shown by the PCV PK/15 strain (Meehan B. et al. Sequence of porcine circovirus DNA; affinities with plant circoviruses, 1997 78, 221-227) which possess in contrast a PstI site and do not possess any EcoRI site.

**Example 9: Cloning of the genome of the porcine circovirus Imp.999 strain**

The restriction fragment of about 1.8 kbp generated by digestion of the double-stranded replicative form of the PCV Imp.999 strain with the restriction enzyme EcoRI was isolated after electrophoresis on a 1.5% agarose gel (see Example 3) using a Qiagen commercial kit (QIAEXII Gel Extraction Kit, Cat # 20021, QIAGEN Ltd., Crawley, West Sussex, UK). This EcoRI-EcoRI restriction fragment was then ligated with the vector pGEM-7 (Promega, Medical Supply Company, Dublin, Ireland), previously digested with the same restriction enzymes and dephosphorylated, according to standard cloning techniques (Sambrook J. et al. Molecular cloning: A Laboratory Manual, 2nd Edition, Cold Spring Harbor Laboratory, Cold Spring Harbor, New York, 1989). The plasmids obtained were

transformed into an *Escherichia coli* JM109 host strain (Stratagene, La Jolla, USA) according to standard techniques. The EcoRI-EcoRI restriction fragment of the PCV Imp.999 strain was also cloned into the EcoRI site of the vector pBlueScript SK+ (Stratagene Inc. La Jolla, USA). Among the clones obtained for each host strain, at least 2 clones containing the fragments of the expected size were selected. The clones obtained were then cultured and the plasmids containing the complete genome of the Imp.999 strain were purified in a small volume (2 ml) or in a large volume (250 ml) according to standard plasmid preparation and purification techniques.

**Example 10: Sequencing of a genomic DNA (double-stranded replicative form) of the PCV Imp.999 strain.**

The nucleotide sequence of 2 EcoRI Imp.999 clones (clones pGEM-7/2 and pGEM-7/8) was determined according to Sanger's dideoxynucleotide technique using the sequencing kit "AmpliTaQ DNA polymerase FS" (Cat # 402079 PE Applied Biosystems, Warrington, UK) and an Applied BioSystems AB1373A automatic sequencing apparatus according to the supplier's recommendations. The initial sequencing reactions were carried out with the M13 "forward" and "reverse" universal primers. The following sequencing reactions were generated according to the "DNA walking" technique. The oligonucleotides necessary for these subsequent sequencings were synthesized by Life Technologies (Inchinnan Business Park, Paisley, UK).

The sequences generated were assembled and analysed by means of the MacDNASIS version 3.2 software (Cat # 22020101, Appligene, Durham, UK). The various open reading frames were analysed by means of the BLAST algorithm available on the "National Center for Biotechnology Information" (NCBI, Bethesda, MD, USA) server.

The complete sequence (EcoRI-EcoRI fragment) is presented in SEQ ID No: 3 (Figure 3). It gives the

total sequence of this strain, which was made to start arbitrarily at the beginning of the EcoRI site, that is to say the G as the first nucleotide.

The procedure was carried out in a similar manner for obtaining the sequence of the other three isolates according to the invention (see SEQ ID No: 1, 2 and 4 and Figures 1, 2 and 4).

The size of the genome of these four strains is:

10	Imp. 1011-48121	1767 nucleotides
	Imp. 1011-48285	1767 nucleotides
	Imp. 999	1768 nucleotides
	Imp. 1010	1768 nucleotides

Example 11: Analysis of the sequence of the PCV Imp.999 strain.

When the sequence generated from the Imp.999 strain was used to test for homology with respect to the sequences contained in the GenBank databank, the only significant homology which was detected is a homology of about 76% (at nucleic acid level) with the sequence of the PK/15 strain (accession numbers Y09921 and U49186) (see Figure No. 5).

At amino acid level, the test for homology in the translation of the sequences in the 6 phases with the databanks (BLAST X algorithm on the NABI server) made it possible to demonstrate a 94% homology with the open reading frame corresponding to the theoretical replicase of the BBTV virus similar to the circoviruses of plants (GenBank identification number 1841515) encoded by the GenBank U49186 sequence.

No other sequence contained in the databanks show significant homology with the sequence generated from the PCV Imp.999 strain.

Analysis of the sequences obtained from the Imp.999 strain cultured using lesions collected from Californian piglets having clinical signs of the multisystemic wasting syndrome shows clearly that this viral isolate is a new porcine circovirus strain.



**Example 12: Comparative analysis of the sequences**

The alignment of the nucleotide sequences of the 4 new PCV strains was made with the sequence of the PCV PK/15 strain (Figure 5). A homology matrix taking into account the four new strains and the previous PK/15 strain was established. The results are the following:

- 1 : Imp. 1011-48121
- 10 2 : Imp. 1011-48285
- 3 : Imp. 999
- 4 : Imp. 1010
- 5: PK/15

	1	2	3	4	5
1	1.0000	0.9977	0.9615	0.9621	0.7600
2		1.0000	0.9621	0.9632	0.7594
3			1.0000	0.9949	0.7560
4				1.0000	0.7566
5					1.0000

The homology between the two French strains Imp. 1011-48121 and Imp. 1011-48285 is greater than 99% (0.9977).

The homology between the two North American strains Imp. 999 and Imp. 1010 is also greater than 99% (0.9949). The homology between the French strains and the North American strains is slightly greater than 96%.

The homology between all these strains and PK/15 falls at a value between 75 and 76%.

It is deduced therefrom that the strains according to the invention are representative of a new type of porcine circovirus, distinct from the type represented by the PK/15 strain. This new type, isolated from pigs exhibiting the PMWS syndrome, is called type II porcine circovirus, PK/15 representing type I. The strains belonging to this type II exhibit remarkable nucleotide sequence homogeneity, although

they have in fact been isolated from very distant geographical regions.

**Example 13: Analysis of the proteins encoded by the genome of the new PCV strains.**

The nucleotide sequence of the Imp. 1010 isolate was considered to be representative of the other circovirus strains associated with the multi-systemic wasting syndrome. This sequence was analysed in greater detail with the aid of the BLASTX algorithm (Altschul et al. J. Mol. Biol. 1990. 215. 403-410) and of a combination of programs from the set of MacVector 6.0 software (Oxford Molecular Group, Oxford OX4 4GA, UK). It was possible to detect 13 open reading frames (or ORFs) of a size greater than 20 amino acids on this sequence (circular genome). These 13 ORFs are the following:

Name	Start	End	Strand	Size of the ORF (nucleotides (nt))	Protein size (amino acids (aa))
ORF1	103	210	sense	108 nt	35 aa
ORF2	1180	1317	sense	138 nt	45 aa
ORF3	1363	1524	sense	162 nt	53 aa
ORF4	398	1342	sense	945 nt	314 aa
ORF5	900	1079	sense	180 nt	59 aa
ORF6	1254	1334	sense	81 nt	26 aa
ORF7	1018	704	antisense	315 nt	104 aa
ORF8	439	311	antisense	129 nt	42 aa
ORF9	190	101	antisense	90 nt	29 aa
ORF10	912	733	antisense	180 nt	59 aa
ORF11	645	565	antisense	81 nt	26 aa
ORF12	1100	1035	antisense	66 nt	21 aa
ORF13	314	1381	antisense	702 nt	213 aa

The positions of the start and end of each ORF refer to the sequence presented in Figure No. 4 (SEQ ID No. 4), of the genome of strain 1010. The

limits of ORFs 1 to 13 are identical for strain 999. They are also identical for strains 1011-48121 and 1011-48285, except for the ORFs 3 and 13:

ORF3 1432-1539, sense, 108 nt, 35aa

5 ORF13 314-1377, antisense, 705 nt, 234 aa.

Among these 13 ORFs, 4 have a significant homology with analogous ORFs situated on the genome of the cloned virus PCV PK-15. Each of the open reading frames present on the genome of all the circovirus  
10 isolates associated with the multisystemic wasting syndrome was analysed. These 4 ORFs are the following:

Name	Start	End	Strand	Size of the ORF (nt)	Protein size (aa)	Molecular mass
ORF4	398	1342	sense	945 nt	314 aa	37.7 kDa
ORF7	1018	704	antisense	315 nt	104 aa	11.8 kDa
ORF10	912	733	antisense	180 nt	59 aa	6.5 kDa
ORF13	314	1381	antisense	702 nt	233 aa	27.8 kDa

15 The positions of the start and end of each ORF refer to the sequence presented in Figure No. 4 (SEQ ID No. 4). The size of the ORF (in nucleotides = nt) includes the stop codon.

20 The comparison between the genomic organization of the PCV Imp. 1010 and PCV PK-15 isolates allowed the identification of 4 ORFs preserved in the genome of the two viruses. The table below presents the degrees of homology observed:

ORF Imp. 1010/ORF PVC PK-15	Percentage homology
ORF4/ORF1	86%
ORF13/ORF2	66.4%
ORF7/ORF3	61.5% (at the level of the overlap (104 aa))
ORF10/ORF4	83% (at the level of the overlap (59 aa))

The greatest sequence identity was observed between ORF4 Imp. 1010 and ORF1 PK-15 (86% homology). This was expected since this protein is probably involved in the replication of the viral DNA and is essential for the viral replication (Meehan et al. J. Gen. Virol. 1997. 78. 221-227; Mankertz et al. J. Gen. Virol. 1998. 79. 381-384).

The sequence identity between ORF13 Imp. 1010 and ORF2 PK-15 is less strong (66.4% homology), but each of these two ORFs indeed exhibits a highly conserved N-terminal basic region which is identical to the N-terminal region of the major structural protein of the CAV avian circovirus (Meehan et al. Arch. Virol. 1992. 124. 301-319). Furthermore, large differences are observed between ORF7 Imp. 1010 and ORF3 PK-15 and between ORF10 Imp. 1010 and ORF4 PK-15. In each case, there is a deletion of the C-terminal region of the ORF7 and ORF10 of the Imp. 1010 isolate when they are compared with ORF3 and ORF4 of PCV PK-15. The greatest sequence homology is observed at the level of the N-terminal regions of ORF7/ORF3 (61.5% homology at the level of the overlap) and of ORF10/ORF4 (83% homology at the level of the overlap).

It appears that the genomic organization of the porcine circovirus is quite complex as a consequence of the extreme compactness of its genome. The major structural protein is probably derived from splicing between several reading frames situated on the same strand of the porcine circovirus genome. It can therefore be considered that any open reading frame (ORF1 to ORF13) as described in the table above can represent all or part of an antigenic protein encoded by the type II porcine circovirus and is therefore potentially an antigen which can be used for specific diagnosis and/or for vaccination. The invention therefore relates to any protein comprising at least one of these ORFs. Preferably, the invention relates to a protein essentially consisting of ORF4, ORF7, ORF10 or ORF13.

**Example 14: Infectious character of the PCV genome cloned from the new strains.**

The plasmid pGEM-7/8 containing the complete  
5 genome (replicative form) of the Imp.999 isolate was  
transfected into PK/15 cells according to the technique  
described by Meehan B. et al. (Characterization of  
viral DNAs from cells infected with chicken anemia  
agent: sequence analysis of the cloned replicative form  
10 and transfection capabilities of cloned genome  
fragments. Arch. Virol. 1992, 124, 301-319).  
Immunofluorescence analysis (see Example 4) carried out  
on the first passage after transfection on  
noncontaminated PK/15 cells have shown that the plasmid  
15 of the clone pGEM7/8 was capable of inducing the  
production of infectious PCV virus. The availability of  
a clone containing an infectious PCV genetic material  
allows any useful manipulation on the viral genome in  
order to produce modified PCV viruses (either  
20 attenuated in pigs, or defective) which can be used for  
the production of attenuated or recombinant vaccines,  
or for the production of antigens for diagnostic kits.

**Example 15: Production of PCV antigens by *in vitro* culture**

The culture of the noncontaminated PK/15 cells  
and the viral multiplication were carried out according  
to the same methods as in Example 1. The infected cells  
are harvested after trypsinization after 4 days of  
30 incubation at 37°C and enumerated. The next passage is  
inoculated with 400,000 infected cells per ml.

**Example 16: Inactivation of the viral antigens**

At the end of the viral culture, the infected  
35 cells are harvested and lysed using ultrasound (Branson  
Sonifier) or with the aid of a rotor-stator type  
colloid mill (UltraTurrax, IKA). The suspension is then  
centrifuged at 3700 g for 30 minutes. The viral  
suspension is inactivated with 0.1% ethyleneimine for

18 hours at +37°C or with 0.5% beta-propiolactone for 24 hours at +28°C. If the virus titre before inactivation is inadequate, the viral suspension is concentrated by ultrafiltration using a membrane with a 300 kDa cut-off (Millipore PTMK300). The inactivated viral suspension is stored at +5°C.

Example 17: Preparation of the vaccine in the form of an emulsion based on mineral oil.

The vaccine is prepared according to the following formula:

- suspension of inactivated porcine circovirus: 250 ml
- Montanide® ISA 70 (SEPPIC): 750 ml

The aqueous phase and the oily phase are sterilized separately by filtration. The emulsion is prepared by mixing and homogenizing the ingredients with the aid of a Silverson turbine emulsifier.

One vaccine dose contains about  $10^{7.5}$  TCID<sub>50</sub>. The volume of one vaccine dose is 0.5 ml for administration by the intradermal route, and 2 ml for administration by the intramuscular route.

This vaccine is used in a vaccination programme against the multisystemic wasting syndrome in combination with the Parvovax® vaccine.

Example 18: Preparation of the vaccine in the form of a metabolizable oil-based emulsion.

The vaccine is prepared according to the following formula:

- suspension of inactivated porcine circovirus: 200 ml
- Dehymuls HRE 7 (Henkel): 60 ml
- Radia 7204 (Oleofina): 740 ml

The aqueous phase and the oily phase are sterilized separately by filtration. The emulsion is prepared by mixing and homogenizing the ingredients with the aid of a Silverson turbine emulsifier.

One vaccine dose contains about  $10^{7.5}$  TCID<sub>50</sub>. The volume of one vaccine dose is 2 ml for administration by the intramuscular route.

This vaccine is used in a vaccination programme against the multisystemic wasting syndrome in combination with the Parvovax® vaccine.

Example 19: The indirect immunofluorescence results in relation to the US and French PCV virus strains and to the PK/15 contaminant with a hyperimmune serum (PCV-T), a panel of monoclonal antibodies F99 prepared from PK/15 and a hyperimmune serum prepared from the Canadian strain (PCV-C)

VIRUS			
	PK/15	USA	France
PCV-T antiserum	≥ 6400	200	800
PCV-C antiserum	200	≥ 6.400	≥ 6.400
F99 1H4	≥ 10 000	<100	100
F99 4B10	≥ 10 000	<100	<100
F99 2B7	≥ 10 000	100	<100
F99 2E12	≥ 10 000	<100	<100
F99 1C9	≥ 10 000	<100	100
F99 2E1	≥ 10 000	<100	<100
F99 1H4	≥ 10 000	100	<100

\* Reciprocal of the last dilution of the serum or of the monoclonal antibody which gives a positive reaction in indirect immunofluorescence.

Example 20: Experimental production of the porcine multisystemic wasting syndrome - protocol 1

Three-day old gnotobiotic piglets obtained by caesarean and kept in an isolating unit were inoculated with virus solutions of PCV. The type II PCV viruses used were the Imp 1010 isolate and the virus obtained from lymph node homogenates obtained from diseased pigs.

Five groups were formed. The piglets were all inoculated at the age of three days by the oronasal route with 1.5 ml of virus solution according to the following scheme:

Group	Number	Virus	Dose
A	6	Lymph node homogenate	ND
B	5	Imp. 1010 (low passage)	10 <sup>2</sup> TCID <sub>50</sub>
C	4	Imp. 1010 (high passage)	10 <sup>2</sup> TCID <sub>50</sub>
D	2	Lysate of PK15 cells free of PCV virus	---
E	3	---	---

#### Results of the experimental challenge:

During the 5-week observation period, the piglets did not develop clinical signs, apart from one animal in group B which showed substantial exhaustion. At autopsy, the pigs in groups A, B and C exhibit hyperplasia of the lymph nodes (size 2 to 10 times greater than that for the animals in groups D and E), in particular of the submaxillary, bronchial, mesenteric, iliac and femoral ganglia. This hyperplasia is linked to a considerable expansion of the cortical zones by infiltration by monocytes and macrophages. The piglets in groups A, B and C also exhibit hyperplasia of the bronchial lymphoid tissue. One piglet in each of groups A, B and C has pneumonia. The piglet in group B, which exhibited substantial exhaustion, and one piglet in group A have a gastric ulcer. Moreover, all the animals in groups A, B and C have myositis in the muscular tunica of the stomach and of the intestine. Most of the animals in groups A, B and C have myocarditis, multifocal hepatitis with lymphocyte, macrophage and eosinophile infiltration, as well as cortical and medullary interstitial nephritis.



One piglet in group C has a liver whose size is bigger than normal, with disseminated clear foci at its surface.

5 No lesion was observed in the piglets in groups D and E.

Circovirus was isolated from the organs of pigs in groups A, B and C.

Example 21: Experimental reproduction of the porcine multisystemic wasting syndrome - protocols 2 and 3

10

Conventional piglets, but isolated from their mother from birth, were inoculated with viral solutions of type II PCV, of porcine parvovirus, or with a mixture of the two viruses.

15 The type II PCV viruses used were the Imp. 1010 and Imp. 1011 isolates (strain 48121).

The PPV virus used is an isolate of Canadian origin, Imp. 1005. This virus has a sequence (1/3 of the sequenced genome) which is identical to that of other  
20 known porcine parvovirus strains (PPV strain NADL-2 and Kresse strain).

Two experimental protocols were carried out.

#### Protocol 2

25 Three groups were formed with 3-day-old piglets. The piglets were all inoculated by the oronasal route with 1 ml of viral solution according to the following scheme:

Group	Number	Virus	Dose
A	5	Imp. 1010	$10^7$ TCID <sub>50</sub>
B	5	Imp. 1010 + Imp. 1005	$5 \times 10^6$ TCID <sub>50</sub>
C (control)	2	---	---

30

Results of the experimental challenge:

Group A: 2 piglets died 21 days after the inoculation and one piglet was humanely killed 24 days after the inoculation.

Group B: 1 piglet died 23 days after the inoculation and one piglet was humanely killed 24 days after the inoculation.

5 The autopsies carried out on the piglets that died following an infection showed the presence of substantial macroscopic lesions: presence of fluid in the pleural cavity, lung oedema, haemorrhages in the kidneys, whitish lesions in the form of a pin head on the kidneys, hepatic necrosis. These lesions are  
10 identical to those observed in the field cases.

The autopsies carried out on the sacrificed piglets did not show macroscopic lesions.

The histological examinations performed on organs removed from the piglets in groups A and B which died  
15 following an infection, as well as in the sacrificed pigs in these 2 groups, showed a typical and complete pattern of the lesions of porcine multisystemic wasting syndrome which are observed in animals in the field:

hepatic necrosis, necrosis of the lymph nodes,  
20 pancreatic necrosis, focal necrosis and severe haemorrhages in the kidneys, presence of syncytia in the lungs, severe necrosis of the hepatocytes with the presence of nuclear inclusions.

It should be noted that a massive quantity of PCV  
25 antigen was found in all these lesions (dead or sacrificed pigs), but that the presence of PPV antigen could not be detected in these same lesions.

No lesion could be detected in the control piglets in group C.

30

### Protocol 3

Four groups were formed with 4-week-old piglets. The pigs were all inoculated by the oronasal route with 1 ml of viral solution according to the following  
35 scheme:

Group	Number	Virus	Dose
A (control)	2	---	---
B	4	Imp. 1005 (PPV)	$10^{5.3}$ TCID50
C	4	Imp. 1011 (PCV)	$10^5$ TCID50
D	4	Imp. 1005 + Imp. 1011	$10^5 + 5 \times 10^4$ TCID50

Results of the experimental challenge:

1 "control" piglet and 2 piglets in each experimental  
5 group (B, C and D) were humanely killed and subjected  
to autopsy 2 weeks after inoculation. Significant  
immunohistological lesions were observed in the two  
piglets in group D (PCV + PPV coinfection). It should  
be noted that it was not possible to detect the  
10 presence of porcine parvovirus in these lesions,  
although a seroconversion in relation to the porcine  
parvovirus was observed in all the pigs in group D.

No macroscopic or histological lesion could be observed  
in the control piglet and in the piglets in the other  
15 groups.

It therefore appears that the PCV + PPV combination  
makes it possible to reproduce histological lesions  
typical of the porcine multisystemic wasting syndrome.  
Following these two experimental protocols, it can be  
20 observed that the inoculation of PCV alone, as a  
PCV + PPV mixture, leads to a more or less severe  
reproduction of the porcine multisystemic wasting  
syndrome, but only the porcine circovirus can be  
detected in the lesions. By contrast, an experimental  
25 infection with PPV alone (group B of protocol 3) does  
not allow macroscopic or histological lesions to be  
induced; however, in the presence of PCV, the  
appearance of lesions is observed in 4-week-old pigs  
(group D of protocol 3).

-32-

## CLAIMS

1. Antigenic preparation directed against the PMWS syndrome, comprising porcine circovirus antigen and  
5 porcine parvovirus antigen.

2. Preparation according to Claim 1, wherein it comprises type II porcine circovirus antigen.

3. Preparation according to claim 2, wherein the  
type II porcine circovirus antigen is an antigen of a  
10 circovirus selected from the group consisting of the  
preparations deposited at the ECACC, under the  
following references :

- accession No. V97100219
- accession No. V97100218
- 15 - accession No. V97100217
- accession No. V98011608
- accession No. V98011609

4. Preparation according to any one of claims 1 to  
3, wherein the porcine circovirus antigen and the  
20 porcine parvovirus antigen comprise, independently of  
each other, an antigen chosen from the group consisting  
of an attenuated live whole antigen, an inactivated  
whole antigen, a subunit antigen, a recombinant live  
vector and a DNA vector.

25 5. Preparation according to anyone of claims 1 to  
4, wherein it comprises, in addition, an other valency  
which corresponds to another pig pathogen.

6. Preparation according to claim 5, wherein it  
comprises an other valency chosen among the group  
30 consisting of : PRRS, Mycoplasma hyopneumoniae,  
Actinobacillus pleuropneumoniae, E. coli, Atrophic  
Rhinitis, Pseudorabies, Hog cholera, Swine Influenza  
and combinations thereof.

-33-

7. Preparation according to claim 5, wherein it comprises an other valency which is PRRS.

8. Vaccine against the PMWS syndrome, comprising an effective quantity of an antigenic preparation  
5 according to any one of claims 1 to 7, in a vehicle or excipient acceptable from the veterinary point of view.

9. Vaccine according to claim 8, wherein it  
10 comprises an adjuvant acceptable from the veterinary point of view.

10. Vaccine according to claim 8 or 9, wherein it comprises antigens of several porcine circoviruses.

11. Vaccine according to any one claims 8 to 10, wherein it comprises circovirus antigen encoded by a  
15 circovirus open reading frame chosen among the group consisting of ORFs 1 to 13.

12. Vaccine according to claim 11, wherein it comprises circovirus antigen encoded by a circovirus open reading frame chosen among the group consisting of  
20 ORFs 4, 7, 10 and 13.

13. Vaccine according to any one of claims 8 to 12, wherein it comprises an expression vector selected from the group consisting of live viruses capable of multiplying in pigs without being pathogenic for pig,  
25 and DNA vectors, this expression vector comprising and expressing said ORF.

14. Vaccine according to claim 13, wherein the viral vector is a virus selected from the group consisting of pig herpes viruses, porcine adenovirus  
30 and poxviruses.

15. Vaccine according to claim 14, wherein the viral vector is a virus selected from the group consisting of Aujeszky's disease virus, vaccinia virus, avipox virus, canarypox virus and swine pox virus.

-34-

16. Vaccination kit containing, packaged separately, a vaccine against the porcine circovirus according to any one of claims 8 to 15, and a vaccine against the porcine parvovirus.

1/ 26

Sequence of the PCV Imp10111-48121 isolate (SEQ ID No. 1)

```
1  AATTCAACCT TAACCTTTCT TATTCTGTAG TATTCAAAGG GCACAGAGCG
51  GGGGTTTGAG CCCCCTCCTG GGGGAAGAAA GTCATTAATA TTGAATCTCA
101 TCATGTCCAC CGCCCAGGAG GCGGTTCTGA CTGTGGTTCG CTTGACAGTA
151 TATCCGAAGG TCGGGGAGAG GCGGGTGTG AAGATGCCAT TTTTCCTTCT
201 CCAGCGGTAA CCGTGGCGGG GGTGGACGAG CCAGGGGCGG CGGCGGAGGA
251 TCTGGCCAAG ATGGCTGCGG GGGCGGTGTC TTCTTCTCCG GTAACGCCTC
301 CTTGGATACG TCATATCTGA AAACGAAAGA AGTGCGCTGT AAGTATTACC
351 AGCGCACTTC GGCAGCGGCA GCACCTCGGC AGCACCTCAG CAGCAACATG
401 CCGAGCAAGA AGAATGGAAG AAGCGGACCC CAACCCCATATA AAGGTGGGT
```

FIG.1

2/26

451 GTTCACTCTG AATAATCCTT CCGAAGACGA GCGCAAGAAA ATACGGGATC  
501 TTCCAATATC CCTATTGAT TATTTATTG TTGGCGAGGA GGGTAATGAG  
551 GAAGGACGAA CACCTCACCT CCAGGGGTC GCTAATTTG TGAAGAAGCA  
601 GACTTTTAAT AAAGTGAAGT GGTATTGGG TGCCCGCTGC CACATCGAGA  
651 AAGCGAAAGG AACAGATCAG CAGAAATAAG AATACTGCAG TAAAGAAGGC  
701 AACTTACTGA TGGAGTGTGG AGCTCCTAGA TCTCAGGGAC AACGGAGTGA  
751 CCTGTCTACT GCTGTGAGTA CCTTGTGGA GAGCGGGAGT CTGGTGACCG  
801 TTGCAGAGCA GCACCCCTGTA ACGTTTGTC GAAATTCCG CGGGCTGGCT  
851 GAACTTTTGA AAGTGAGCGG GAAAATGCAG AAGCGTGATT GGAAGACTAA  
901 TGTacACGTC ATTGTGGGC CACCTGGGTG TGGTAAAGC AAATGGGCTG

FIG.1 (cont.)



3/ 26

951 CTAATTTTGC AGACCCGGAA ACCACATACT GGAACCACC TAGAAACAAG  
1001 TGGTGGGATG GTTACCATGG TGAAGAAGTG GTTGTATTG ATGACTTTTA  
1051 TGGCTGGCTG CCCTGGGATG ATCTACTGAG ACTGTGTGAT CGATATCCAT  
1101 TGA CTGTAGA GACTAAAGGT GGA ACTGTAC CTTTTTTGGC CCGCAGTATT  
1151 CTGATTACCA GCAATCAGAC CCCGTTGGAA TGGTACTCCT CAACTGCTGT  
1201 CCCAGCTGTA GAAGCTCTTT ATCGGAGGAT TACTTCCTTG GTATTTTGGA  
1251 AGAATGCTAC AGAACAATCC ACGGAGGAAG GGGGCCAGTT CGTCACCCCTT  
1301 TCCCCCCCAT GCCCTGAATT TCCATATGAA ATAAATTACT GAGTCTTTTT  
1351 TATCACTTCG TAATGGTTTT TATTATTCAT TAAGGGTTAA GTGGGGGGTC  
1401 TTTAAGATTA AATCTCTGA ATTGTACATA CATGGTTACA CGGATATTGT

FIG.1 (cont.)

4/26

1451 ATTCCTGGTC GTATATACTG TTTTCGAACG CAGTGCCGAG GCCTACGTGG  
1501 TCACATTTC CAGCAGTTTG TAGTCTCAGC CACAGCTGGT TTCTTTTGT  
1551 GTTGGTTGG AAGTAATCAA TAGTGGAATC TAGGACAGGT TTGGGGGTAA  
1601 AGTAGCGGGA GTGGTAGGAG AAGGCTGGG TTATGGTATG GCGGAGGAG  
1651 TAGTTTACAT AGGGTCATA GGTAGGGCT GTGGCCTTG TTACAAAGTT  
1701 ATCATCTAGA ATAACAGCAC TGGAGCCAC TCCCCTGTCA CCTGGGTGA  
1751 TCGGGGAGCA GGGCCAG

FIG.1 (cont. and end)

5/26

Sequence of the PCV Imp1011-48285 isolate (SEQ ID No. 2)

```
1  AATTCAACCT TAACCTTTCT TATTCTGTAG TATTCAAAGG GCACAGAGCG
51  GGGGTTTGAG CCCCCTCCTG GGGGAAGAAA GTCATTAATA TTGAATCTCA
101 TCATGTCCAC CGCCCAGGAG GCGGTTTGA CTGTGGTTCC CTTGACAGTA
151 TATCCGAAGG TCGGGGAGAG GCGGGTGTG AAGATGCCAT TTTTCCTTCT
201 CCAGCGGTAA CCGTGCGCGG GGTGGACGAG CCAGGGGCGG CGGCGGAGGA
251 TCTGGCCAAG ATGGCTGCGG GGGCGGTGTC TTCTTCTCCG GTAACGCCCTC
301 CTTGGATACG TCATATCTGA AAACGAAAGA AGTCCGCTGT AAGTATTACC
351 AGCGCACTC GGCAGCGGCA GCACCTCGGC AGCACCTCAG CAGCAACATG
401 CCCAGCAAGA AGAATGGAAG AAGCGGACCC CAACCCCAT AAGGTGGGT
451 GTTCACTCTG AATAATCCTT CCGAAGACGA GCGCAAGAAA ATACGGGATC
```

FIG. 2

6 / 26

501 TTCCAATATC CCTATTGAT TATTTATTG TTGCGAGGA GGTAAATGAG  
551 GAAGGACGAA CACCTCACCT CCAGGGGTTT GCTAATTTTG TGAAGAAGCA  
601 GACTTTTAAT AAAGTGAAGT GGTATTGGG TGCCCGCTGC CACATCGAGA  
651 AAGCGAAAGG AACAGATCAG CAGAATAAAG AATACTGCAG TAAAGAAGGC  
701 AACTTACTGA TGGAGTGTGG AGCTCCTAGA TCTCAGGGAC AACGGAGTGA  
751 CCTGTCTACT GCTGTGAGTA CCTTGTTGGA GAGCGGGAGT CTGGTGACCG  
801 TTGCAGAGCA GCACCCCTGTA ACGTTTGTCA GAAATTTCG CGGGCTGGCT  
851 GAACTTTTGA AAGTGAGCGG GAAATGCAG AAGCGTGATT GGAAGACTAA  
901 TGTACACGTC ATTGTGGGGC CACCTGGGTG TGGTAAAGC AAATGGGCTG  
951 CTAATTTTGC AGACCCGGAA ACCACATACT GGAAACCACC TAGAAACAAG  
1001 TGGTGGGATG GTTACCATGG TGAAGAAGTG GTTGTATTG ATGACTTTTA

FIG. 2 (cont.)

7/ 26

1051	TGGCTGGCTG	CCCTGGGATG	ATCTACTGAG	ACTGTGTGAT	CGATATCCAT
1101	TGACTGTAGA	GACTAAAGGT	GGAAGTGTAC	CTTTTGTGGC	CCGCAGTATT
1151	CTGATTACCA	GCAATCAGAC	CCCGTTGGAA	TGGTACTCCT	CAACTGCTGT
1201	CCCAGCTGTA	GAAGCTCTTT	ATCGGAGGAT	TACTTCCTTG	GTATTTTGGA
1251	AGAATGCTAC	AGAACAATCC	ACGAGGAAG	GGGGCCAGTT	CGTCACCCCTT
1301	TCCCCCCCAT	GCCCTGAATT	TCCATATGAA	ATAAATTACT	GAGTCTTTT
1351	TATCACTTCG	TAATGGTTTT	TATTATTCA	TAAGGGTTAA	GTGGGGGGTC
1401	TTTAAGATTA	AATTCTCTGA	ATTGTACATA	CATGGTTACA	CGGATATTGT
1451	ATTCCTGGTC	GTATATACTG	TTTTCGAACG	CAGTGCCGAG	GCCTACGTGG
1501	TCTACATTTC	CAGTAGTTTG	TAGTCTCAGC	CACAGCTGAT	TTCTTTTGT

FIG.2 (cont.)

8 / 26

1551 GTTTGGTTGG AAGTAATCAA TAGTGAATC TAGGACAGGT TTGGGGGTAA  
1601 AGTAGCGGGA GTGGTAGGAG AAGGGCTGGG TTATGGTATG GCGGGAGGAG  
1651 TAGTTTACAT AGGGGTCATA GGTGAGGGCT GTGGCCCTTG TTACAAAGTT  
1701 ATCATCTAGA ATAAACAGCAC TGGAGCCCAC TCCCCTGTCA CCCTGGGTGA  
1751 TCGGGGAGCA GGGCCAG

FIG.2 (cont. and end)

9/26

Sequence of the PCV Imp999 isolate (SEQ ID No. 3)

```
1  AATTCAACCT TAACCTTTTT TATTCTGTAG TATTCAAAGG GTATAGAGAT
51  TTTGTTGGTC CCCCCTCCCG GGGGAACAAA GTCGTCAATA TTAAATCTCA
101 TCATGTCCAC CGCCCAGGAG GCGGTTCTGA CTGTGGTAGC CTTGACAGTA
151 TATCCGAAGG TCGGGGAGAG GCGGGTGTG AAGATGCCAT TTTTCCCTTCT
201 CCAACGGTAG CGGTGGCGGG GGTGGACGAG CCAGGGGCGG CGGCGGAGGA
251 TCTGGCCAAG ATGGCTGCGG GGGCGGTGTC TTCTTCTGCG GTAACGCCTC
301 CTTGGATACG TCATAGCTGA AAACGAAAGA AGTGCCTGT AAGTATTACC
351 AGCGCACTTC GGCAGCGGCA GCACCTCGGC AGCACCTCAG CAGCAACATG
401 CCCAGCAAGA AGAATGGAAG AAGCGGACCC CAACCACATA AAAGGTGGGT
```

FIG.3

10/ 26

451 GTTCACGCTG AATAATCCTT CCGAAGACGA GCGCAAGAAA ATACGGGAGC  
501 TCCCAATCTC CCTATTGAT TATTTATTG TTGGCGAGGA GGGTAATGAG  
551 GAAGGACGAA CACCTCACCT CCAGGGGTTT GCTAATTTTG TGAAGAGGCA  
601 AACTTTTAAT AAAGTGAAGT GGTATTGGG TGCCCGCTGC CACATCGAGA  
651 AAGCCAAAGG AACTGATCAG CAGAATAAG AATATTGCAG TAAAGAAGGC  
701 AACTTACTTA TTGAATGTGG AGTCCTCTCGA TCTCAAGGAC AACGGAGTGA  
751 CCTGTCTACT GCTGTGAGTA CCTTGTGGA GAGCGGGAGT CTGGTGACCG  
801 TTGCAGAGCA GCACCCTGTA ACGTTTGTC GAAATTTCCG CGGGCTGGCT  
851 GAACTTTTGA AAGTGAGCGG GAAATGCAG AAGCGTGATT GGAAGACCAA  
901 TGTACACGTC ATTGTGGGC CACCTGGGTG TGTAAAGC AAATGGGCTG  
951 CTAATTTTGC AGACCCGGA ACCACATACT GGAAACCACC TAGAAACAAG

FIG.3 (cont.)



11/ 26

1001 TGGTGGGATG GTTACCATGG TGAAGAAAGTG GTTGTATTG ATGACTTTTA  
1051 TGGCTGGCTG CCGTGGGATG ATCTACTGAG ACTGTGTGAT CGATATCCAT  
1101 TGA CTGTAGTA GACTAAAGGT GGA ACTGTAC CTTTTTTGGC CCGCAGTATT  
1151 CTGATTACCA GCAATCAGAC CCCGTTGGAA TGGTACTCCT CAACTGCTGT  
1201 CCCAGCTGTA GAAGCTCTCT ATCGGAGGAT TACTTCCTTG GTATTTTGGGA  
1251 AGAATGCTAC AGAACAAATCC ACGGAGGAAG GGGGCCAGTT CGTCACCCCTT  
1301 TCCCCCCCCAT GCCCTGAATT TCCATATGAA ATAAATTACT GAGTCTTTTT  
1351 TATCACTTCG TAATGGTTTT TATTATTCAT TTAGGGTTTA AGTGGGGGGT  
1401 CTTTAAGATT AAATTCTCTG AATTGTACAT ACATGGTTAC ACGGATATTG  
1451 TAGTCCTGGT CGTATATACT GTTTTCGAAC GCAGTGCCGA GGCCTACGTG

FIG.3 (cont.)

12/ 26

1501 GTCCACATTT CTAGAGGTTT GTAGCCTCAG CCAAGCTGA TTCCTTTTGT  
1551 TATTGGTTG GAAGTAATCA ATAGTGGAGT CAAGAACAGG TTTGGGTGTG  
1601 AAGTAACGGG AGTGGTAGGA GAAGGTTGG GGGATTGTAT GCGGGAGGA  
1651 GTAGTTTACA TATGGGTCAT AGTTAGGGC TGTGGCCTTT GTTACAAAGT  
1701 TATCATCTAG AATAACAGCA GTGGAGCCCA CTCCTCTATC ACCCTGGGTG  
1751 ATGGGGGAGC AGGGCCAG

FIG.3 (cont. and end)

13/26

Sequence of the PCV Imp1010 isolate (SEQ ID No. 4)

```
1  AATTCAACCT FAACCTTTCT TATTCTGTAG TATTCAAAGG GTATAGAGAT
51  TTTGTTGGTC CCCCTCCCG GGGAAACAAA GTCGTCAATT TTAAATCTCA
101 TCATGTCCAC CGCCAGGAG GCGTTGTGA CTGTGGTACG CTGACAGTA
151 TATCCGAAGG TCGGGGAGAG GCGGTGTG AAGATGCCAT TTTTCCTTCT
201 CCAACGGTAG CGGTGGCGG GGTGGACGAG CCAGGGGCGG CGCGGAGGA
251 TCTGGCCAAG ATGGCTGCGG GGGCGGTGC TTCTTCTGCG GTAACGCCCTC
301 CTTGGATACG TCATAGCTGA AAACGAAAGA AGTGCCTGT AAGTATTACC
351 AGCGCACTTC GGCAGCGGCA GCACCTCGG AGCACCTCAG CAGCAACATG
401 CCCAGCAAGA AGAATGGAAG AAGCGGACCC CAACCACATA AAAGGTGGGT
451 GTTCACGCTG AATAATCCTT CCGAAGACGA GCGCAAGAAA ATACGGGAGC
```

FIG.4

14/26

501 TCCCAAATCTC CCTATTGAT TATTTTATTG TTGGCGAGGA GGGTAATGAG  
551 GAAGGACGAA CACCTCACCT CCAGGGGTC GCTAATTTTG TGAAGAAGCA  
601 AACTTTTAAT AAAGTGAAGT GGTATTGGG TGCCCGCTGC CACATCGAGA  
651 AAGCCAAAGG AACTGATCAG CAGAAATAAG AATATTGCAG TAAAGAAGGC  
701 AACTTACTTA TTGAATGTGG AGCTCCTCGA TCTCAAGGAC AACGGAGTGA  
751 CCTGTCTACT GCTGTGAGTA CCTTGTGTGA GAGCGGGAGT CTGGTGACCG  
801 TTGCAGAGCA GCACCCTGTA ACGTTTGTCA GAAATTTCG CGGGCTGGCT  
851 GAACTTTTGA AAGTGAGCGG GAAATGCAG AAGCGTGATT GGAAGACCAA  
901 TGTACACGTC ATTGTGGGC CACCTGGGTG TGGTAAAGC AAATGGGCTG  
951 CTAATTTTGC AGACCCGGAA ACCACATACT GGAAACCACC TAGAAACAAG  
1001 TGGTGGGATG GTTACCATGG TGAAGAAGTG GTTGTATTG ATGACTTTTA

FIG.4 (cont.)

15/26

1051 TGGCTGGCTG CCGTGGGATG ATCTACTGAG ACTGTGTGAT CGATATCCAT  
1101 TGA CTGTAGA GACTAAAGGT GGA ACTGTAC CTTTTTTGGC CCGCAGTATT  
1151 CTGATTACCA GCAATCAGAC CCCGTTGGAA TGGTACTCCT CAACTGCTGT  
1201 CCCAGCTGTA GAAGCTCTCT ATCGGAGGAT TACTTCCTTG GTATTTTGGG  
1251 AGAATGCTAC AGAACAATCC ACGGAGGAAG GGGGCCAGTT CGTCACCCTT  
1301 TCCCCCCCCAT GCCCTGAATT TCCATATGAA ATAAATTACT GAGTC TTTT  
1351 TATCACTTCG TAATGGTTTT TATTATTCAT TTAGGGTTTA AGTGGGGGGT  
1401 CTTTAAGATT AAATCTCTCG AATTGTACAT ACATGGTTAC ACGGATATTG  
1451 TAGTCCTGGT CGTATT TACT GTTTTCGAAC GCAGCGCCGA GGCCTACGTG

FIG.4 (cont.)

16/26

1501 GTCCACATTT CCAGAGGTTT GTAGTCTCAG CCAAGCTGA TTCCTTTGT  
1551 TATTTGGTTG GAAGTAATCA ATAGTGGAGT CAAGAACAGG TTTGGGTGTG  
1601 AAGTAACGGG AGTGGTAGGA GAAGGTTGG GGGATTGTAT GCGGGAGGA  
1651 GTAGTTTACA TATGGGTCAT AGGTTAGGC TGTGGCCTTT GTTACAAAGT  
1701 TATCATCTAG AATAACAGCA GTGGAGCCA CTCCCCTATC ACCCTGGGTG  
1751 ATGGGGGAGC AGGCCAG

FIG.4 (cont. and end)

17/26

## CLUSTAL W multiple sequence alignment

```

PCVPK-15      AATTCATATTAGCCTTCTAATACGGTAGTATTGGAAGGTAGGGGTAGGGGGTTGGTG
IMP999-ECO    AATTCAACCTTAACCTTTTATTCTGTAGTATTCAAAGGGTATAGAGATTTTGTGTC
IMP1010-ST    AATTCAACCTTAACCTTTCTTATTCTGTAGTATTCAAAGGGTATAGAGATTTTGTGTC
IMP1011-48    AATTCAACCTTAACCTTTCTTATTCTGTAGTATTCAAAGGGCACAGAGCGGGGTTTGAG
IMP1011-48    AATTCAACCTTAACCTTTCTTATTCTGTAGTATTCAAAGGGCACAGAGCGGGGTTTGAG
                **** * * * * * * * * * * * * * * * *
PCVPK-15      CCGCCTGAGGGGGGAGGAAGTGGCCGATGTTGAATTTGAGGTAGTTAACATTCCAAGAT
IMP999-ECO    CCCCCTCCCGGGGAACAAAGTCGTCAATATTAAATCTCATGTCCACCGCCAGGAG
IMP1010-ST    CCCCCTCCCGGGGAACAAAGTCGTCAATTTTAAATCTCATGTCCACCGCCAGGAG
IMP1011-48    CCCCCTCCTGGGGAAGAAAGTCATTAATAATTGAATCTCATGTCCACCGCCAGGAG
IMP1011-48    CCCCCTCCTGGGGAAGAAAGTCATTAATAATTGAATCTCATGTCCACCGCCAGGAG
                ** ** * * * * * * * * * * * * * * * *
PCVPK-15      GGC--TGCGAGTATCCTCCTTTT-ATGGTGAGTACAAATTCTGTAGAAAGCGGGAATTG
IMP999-ECO    GCGTTCTGACTGTGGTAGCCTTGACAGTATATCCGAAGGTGCGGAGAGCGGGTGTG
IMP1010-ST    GCGTTGTGACTGTGGTACGCTTGACAGTATATCCGAAGGTGCGGAGAGCGGGTGTG
IMP1011-48    GCGTTCTGACTGTGGTTCGCTTGACAGTATATCCGAAGGTGCGGAGAGCGGGTGTG
IMP1011-48    GCGTTTGTGACTGTGGTTCGCTTGACAGTATATCCGAAGGTGCGGAGAGCGGGTGTG
                *** * * * * * * * * * * * * * * * *

```

FIG.5

18/26

PCVPK-15 AAGATACCCGTCTTTCGGCGCCATCTGTAAAGGTTTCTGAAGCGGGGTGTGCCAATAT  
 IMP999-ECO AAGATGCCATTTTTCCTTCTCCAACGGTAGCGGTGGC-GGGGTGGA-CGAGCCAGGGGC  
 IMP1010-ST AAGATGCCATTTTTCCTTCTCCAACGGTAGCGGTGGC-GGGGTGGA-CGAGCCAGGGGC  
 IMP1011-48 AAGATGCCATTTTTCCTTCTCCAGCGGTAAACGGTGGC-GGGGTGGA-CGAGCCAGGGGC  
 IMP1011-48 AAGATGCCATTTTTCCTTCTCCAGCGGTAAACGGTGGC-GGGGTGGA-CGAGCCAGGGGC  
 \*\*\*\*\* \* \* \* \* \*

PCVPK-15 GGTCTTCTCCGGAGGATGTTTCCAAGATGGCTCGGGGCGGGTCTTCTTCTGCGGTAA  
 IMP999-ECO GG- - - CGGCGGAGGATCTGGCCAAAGATGGCTCGGGGCGGGTCTTCTTCTGCGGTAA  
 IMP1010-ST GG- - - CGGCGGAGGATCTGGCCAAAGATGGCTCGGGGCGGGTCTTCTTCTGCGGTAA  
 IMP1011-48 GG- - - CGGCGGAGGATCTGGCCAAAGATGGCTCGGGGCGGGTCTTCTTCTGCGGTAA  
 IMP1011-48 GG- - - CGGCGGAGGATCTGGCCAAAGATGGCTCGGGGCGGGTCTTCTTCTGCGGTAA  
 \*

PCVPK-15 CGCCTCCTTGCCACGTCACTCTATAAAGTGAAAGAGTGGCTGCTGTAGTATTACCA  
 IMP999-ECO CGCCTCCTTGGTACGTCTAGC-TGAAACGAAAGAGTGGCTGTA--AGTATTACCA  
 IMP1010-ST CGCCTCCTTGGTACGTCTAGC-TGAAACGAAAGAGTGGCTGTA--AGTATTACCA  
 IMP1011-48 CGCCTCCTTGGTACGTCTATC-TGAAACGAAAGAGTGGCTGTA--AGTATTACCA  
 IMP1011-48 CGCCTCCTTGGTACGTCTATC-TGAAACGAAAGAGTGGCTGTA--AGTATTACCA  
 \*

FIG.5 (cont.)



PCVPK-15  
IMP999-ECO  
IMP1010-ST  
IMP1011-48  
IMP1011-48

PCVPK-15  
IMP999-ECO  
IMP1010-ST  
IMP1011-48  
IMP1011-48

PCVPK-15  
IMP999-ECO  
IMP1010-ST  
IMP1011-48  
IMP1011-48

FIG. 5 (cont.)

PCVPK-15  
 IMP999-ECO  
 IMP1010-ST  
 IMP1011-48  
 IMP1011-48  
 CCGAGAGGAAGGTTTGAAGAGGGTAGAACTCCTCACCTCCAGGGGTTTGGAAATTTTGC  
 TGGCGAGGAGGGTAATGAGGAAGGACGAACACCTCACCTCCAGGGGTTGCTAAATTTTGT  
 TGGCGAGGAGGGTAATGAGGAAGGACGAACACCTCACCTCCAGGGGTTGCTAAATTTTGT  
 TGGCGAGGAGGGTAATGAGGAAGGACGAACACCTCACCTCCAGGGGTTGCTAAATTTTGT  
 \*\* \*\*\*\* \*\* \*\* \*\* \*\* \*\*\*\* \* \*\*\*\* \* \*\*\*\* \* \*\*\*\* \* \*\*\*\* \*  
 TAAGAAGCAGACTTTTAAACAAGGTGAAGTGGTATTTGGTGCCCGCTGCCACATCGAGAA  
 GAAGAAGCAAACTTTTAAATAAAGTGAAGTGGTATTTGGTGCCCGCTGCCACATCGAGAA  
 GAAGAAGCAAACTTTTAAATAAAGTGAAGTGGTATTTGGTGCCCGCTGCCACATCGAGAA  
 GAAGAAGCAGACTTTTAAATAAAGTGAAGTGGTATTTGGTGCCCGCTGCCACATCGAGAA  
 \*\*\*\*\* \*\*\*\* \* \* \*\*\*\* \* \*\*\*\* \* \*\*\*\* \* \*\*\*\* \* \*\*\*\* \*  
 AGCGAAAGGAACCGACCAGCAGATAAAGAACTACTGCAGTAAAGGCCACATACTTAT  
 AGCCAAAGGAAGTATCAGCAGATAAAGAACTATTGCAGTAAAGGCCAATTACTTAT  
 AGCCAAAGGAAGTATCAGCAGATAAAGAACTATTGCAGTAAAGGCCAATTACTTAT  
 AGCGAAAGGAACAGATCAGCAGATAAAGAACTACTGCAGTAAAGGCCAATTACTTAT  
 \*\* \*\*\*\* \* \* \*\*\*\* \* \*\*\*\* \* \*\*\*\* \* \*\*\*\* \* \*\*\*\* \* \*\*\*\* \*  
 PCVPK-15  
 IMP999-ECO  
 IMP1010-ST  
 IMP1011-48  
 IMP1011-48  
 CCGAGAGGAGGTTTGAAGAGGGTAGAACTCCTCACCTCCAGGGGTTTGGAAATTTTGC  
 TGGCGAGGAGGGTAATGAGGAAGGACGAACACCTCACCTCCAGGGGTTGCTAAATTTTGT  
 TGGCGAGGAGGGTAATGAGGAAGGACGAACACCTCACCTCCAGGGGTTGCTAAATTTTGT  
 TGGCGAGGAGGGTAATGAGGAAGGACGAACACCTCACCTCCAGGGGTTGCTAAATTTTGT  
 \*\* \*\*\*\* \* \* \*\*\*\* \* \*\*\*\* \* \*\*\*\* \* \*\*\*\* \* \*\*\*\* \* \*\*\*\* \*  
 TAAGAAGCAGACTTTTAAACAAGGTGAAGTGGTATTTGGTGCCCGCTGCCACATCGAGAA  
 GAAGAAGCAAACTTTTAAATAAAGTGAAGTGGTATTTGGTGCCCGCTGCCACATCGAGAA  
 GAAGAAGCAAACTTTTAAATAAAGTGAAGTGGTATTTGGTGCCCGCTGCCACATCGAGAA  
 GAAGAAGCAGACTTTTAAATAAAGTGAAGTGGTATTTGGTGCCCGCTGCCACATCGAGAA  
 \*\*\*\*\* \*\*\*\* \* \* \*\*\*\* \* \*\*\*\* \* \*\*\*\* \* \*\*\*\* \* \*\*\*\* \*  
 AGCGAAAGGAACCGACCAGCAGATAAAGAACTACTGCAGTAAAGGCCACATACTTAT  
 AGCCAAAGGAAGTATCAGCAGATAAAGAACTATTGCAGTAAAGGCCAATTACTTAT  
 AGCCAAAGGAAGTATCAGCAGATAAAGAACTATTGCAGTAAAGGCCAATTACTTAT  
 AGCGAAAGGAACAGATCAGCAGATAAAGAACTACTGCAGTAAAGGCCAATTACTTAT  
 \*\* \*\*\*\* \* \* \*\*\*\* \* \*\*\*\* \* \*\*\*\* \* \*\*\*\* \* \*\*\*\* \* \*\*\*\* \*

FIG.5 (cont.)

PCVPK-15  
 IMP999-ECO  
 IMP1010-ST  
 IMP1011-48  
 IMP1011-48

GCGCACTTCGGCAGCGGCAGCACCTCGGCAGCG--TCAGTG--AAAATGCCAAGCAAGAA  
 GCGCACTTCGGCAGCGGCAGCACCTCGGCAGCACCTCAGCAGCAACATGCCCAGCAAGAA  
 GCGCACTTCGGCAGCGGCAGCACCTCGGCAGCACCTCAGCAGCAACATGCCCAGCAAGAA  
 GCGCACTTCGGCAGCGGCAGCACCTCGGCAGCACCTCAGCAGCAACATGCCGAGCAAGAA  
 GCGCACTTCGGCAGCGGCAGCACCTCGGCAGCACCTCAGCAGCAACATGCCCAGCAAGAA  
 \*\*\*\*\*

PCVPK-15  
 IMP999-ECO  
 IMP1010-ST  
 IMP1011-48  
 IMP1011-48

PCVPK-15  
 IMP999-ECO  
 IMP1010-ST  
 IMP1011-48  
 IMP1011-48

-----AAGCGGCCCGCAACCCCATAGAGGTGGGTGTTCAACCCTTAATAATCCTTC  
 GAATGGAAGAGCGGACCCCAACACATATAAAGGTGGGTGTTCAACGCTGAATAATCCTTC  
 GAATGGAAGAGCGGACCCCAACACATATAAAGGTGGGTGTTCAACGCTGAATAATCCTTC  
 GAATGGAAGAGCGGACCCCAACACATATAAAGGTGGGTGTTCACTCTGAATAATCCTTC  
 GAATGGAAGAGCGGACCCCAACCCCATATAAAGGTGGGTGTTCACTCTGAATAATCCTTC  
 \*\*\*\*\*

PCVPK-15  
 IMP999-ECO  
 IMP1010-ST  
 IMP1011-48  
 IMP1011-48

CGAGGAGGAGAAACAAAATACGGGAGCTTCCAATCTCCCTTTTGGATTATTTGTTTG  
 CGAAGACGAGCGCAAGAAAATACGGGAGCTCCCAATCTCCCTATTTGATTATTTATTTGT  
 CGAAGACGAGCGCAAGAAAATACGGGAGCTCCCAATCTCCCTATTTGATTATTTATTTGT  
 CGAAGACGAGCGCAAGAAAATACGGGATCTTCCAATATCCCTATTTGATTATTTATTTGT  
 CGAAGACGAGCGCAAGAAAATACGGGATCTTCCAATATCCCTATTTGATTATTTATTTGT  
 \*\*\*\*\*

PCVPK-15  
 IMP999-ECO  
 IMP1010-ST  
 IMP1011-48  
 IMP1011-48

FIG. 5 (cont.)

PCVPK-15

IMP999-ECO

IMP1010-ST

IMP1011-48

IMP1011-48

CGGAGAGGAAGTTTGAAGAGGGTAGAACTCCTCACCTCCAGGGTTTGGAAATTTTGC  
TGGCGAGGAGGGTAATGAGGAAGGACGAACACCTCACCTCCAGGGTTGCGTAATTTTGT  
TGGCGAGGAGGGTAATGAGGAAGGACGAACACCTCACCTCCAGGGTTGCGTAATTTTGT  
TGGCGAGGAGGGTAATGAGGAAGGACGAACACCTCACCTCCAGGGTTGCGTAATTTTGT  
TGGCGAGGAGGGTAATGAGGAAGGACGAACACCTCACCTCCAGGGTTGCGTAATTTTGT  
\*\* \*\*\*\*\* \*\* \*\* \*\* \*\*\*\*\* \*\* \*\*\*\*\*

PCVPK-15

IMP999-ECO

IMP1010-ST

IMP1011-48

IMP1011-48

TAAGAAGCAGACTTTTAACAAGTGAAAGTGGTATTTTGGTGCCCGCTGCCACATCGAGAA  
GAAGAAGCAAACTTTTAATAAAGTGAAAGTGGTATTTGGGTGCCCCGCTGCCACATCGAGAA  
GAAGAAGCAAACTTTTAATAAAGTGAAAGTGGTATTTGGGTGCCCCGCTGCCACATCGAGAA  
GAAGAAGCAGACTTTTAATAAAGTGAAAGTGGTATTTGGGTGCCCCGCTGCCACATCGAGAA  
GAAGAAGCAGACTTTTAATAAAGTGAAAGTGGTATTTGGGTGCCCCGCTGCCACATCGAGAA  
\*\*\*\*\* \*\* \*\*\*\*\* \*\* \*\*\*\*\*

PCVPK-15

IMP999-ECO

IMP1010-ST

IMP1011-48

IMP1011-48

AGCGAAAGGAACCGACCAGCAGAAATAAAGAAATACTGCAGTAAAGAGGCCACATATTAT  
AGCCAAAGGAACCTGATCAGCAGAAATAAAGAAATAATTCAGTAAAGAGGCAACTTACTTAT  
AGCCAAAGGAACCTGATCAGCAGAAATAAAGAAATAATTCAGTAAAGAGGCAACTTACTTAT  
AGCGAAAGGAACAGATCAGCAGAAATAAAGAAATACTGCAGTAAAGAGGCAACTTACTGAT  
AGCGAAAGGAACAGATCAGCAGAAATAAAGAAATACTGCAGTAAAGAGGCAACTTACTGAT  
\*\*\*\*\* \*\* \*\*\*\*\* \*\* \*\*\*\*\*

FIG.5 (cont.)

PCVPK-15  
IMP999-ECO  
IMP1010-ST  
IMP1011-48  
IMP1011-48

CGAGTGTGAGCTCCGCGGAACAGGGGAAGCGAGCACCTGTCTACTGCTGTGAGTAC  
TGAATGTGAGCTCCTCGATCTCAAGGACAACGGAGTGACCTGTCTACTGCTGTGAGTAC  
TGAATGTGAGCTCCTCGATCTCAAGGACAACGGAGTGACCTGTCTACTGCTGTGAGTAC  
GGAGTGTGAGCTCCTAGATCTCAGGGACAACGGAGTGACCTGTCTACTGCTGTGAGTAC  
GGAGTGTGAGCTCCTAGATCTCAGGGACAACGGAGTGACCTGTCTACTGCTGTGAGTAC  
\* \* \* \* \*

PCVPK-15  
IMP999-ECO  
IMP1010-ST  
IMP1011-48  
IMP1011-48

CCTTTTGGAGACGGGTCCTTTGGTGACTGTAGCCGAGCAGTTCCCTGTAACGTATGTGAG  
CTTGTGTGAGACGGGAGTCTGGTGACCGTTGCAGAGCAGCACCCCTGTAACGTTTGTGAG  
CTTGTGTGAGACGGGAGTCTGGTGACCGTTGCAGAGCAGCACCCCTGTAACGTTTGTGAG  
CTTGTGTGAGACGGGAGTCTGGTGACCGTTGCAGAGCAGCACCCCTGTAACGTTTGTGAG  
CTTGTGTGAGACGGGAGTCTGGTGACCGTTGCAGAGCAGCACCCCTGTAACGTTTGTGAG  
\* \* \* \* \*

PCVPK-15  
IMP999-ECO  
IMP1010-ST  
IMP1011-48  
IMP1011-48

AAATTTCCGCGGCTGGCTGAACCTTTTGAAGTGAGCGGGGAAGATGCAGCAGCGTGATTG  
AAATTTCCGCGGCTGGCTGAACCTTTTGAAGTGAGCGGGGAAGATGCAGCAGCGTGATTG  
AAATTTCCGCGGCTGGCTGAACCTTTTGAAGTGAGCGGGGAAGATGCAGCAGCGTGATTG  
AAATTTCCGCGGCTGGCTGAACCTTTTGAAGTGAGCGGGGAAGATGCAGCAGCGTGATTG  
AAATTTCCGCGGCTGGCTGAACCTTTTGAAGTGAGCGGGGAAGATGCAGCAGCGTGATTG  
\* \* \* \* \*

FIG.5 (cont.)

22/26

PCVPK-15  
IMP999-ECO  
IMP1010-ST  
IMP1011-48  
IMP1011-48

GAAGACAGCTGTACACGTCAATAGTGGCCCGCCCGTGTGTGGAAAGAGCCAGTGGCCCG  
GAAGACCAATGTACACGTCAATGTGGGGCCACCTGGGTGTGTTAAAGCAAAATGGGCTGC  
GAAGACCAATGTACACGTCAATGTGGGGCCACCTGGGTGTGTTAAAGCAAAATGGGCTGC  
GAAGACTAATGTACACGTCAATGTGGGGCCACCTGGGTGTGTTAAAGCAAAATGGGCTGC  
GAAGACTAATGTACACGTCAATGTGGGGCCACCTGGGTGTGTTAAAGCAAAATGGGCTGC  
\*\*\*\*\*

PCVPK-15  
IMP999-ECO  
IMP1010-ST  
IMP1011-48  
IMP1011-48

TAAATTTGCTGAGCCTAGGGACACCTACTGGAAGCCTAGTAGAAATAAGTGTGGATGG  
TAAATTTGCAGACCCGGAACCCACATACTGGAACCCACCTAGAAAACAAGTGTGGATGG  
TAAATTTGCAGACCCGGAACCCACATACTGGAACCCACCTAGAAAACAAGTGTGGATGG  
TAAATTTGCAGACCCGGAACCCACATACTGGAACCCACCTAGAAAACAAGTGTGGATGG  
TAAATTTGCAGACCCGGAACCCACATACTGGAACCCACCTAGAAAACAAGTGTGGATGG  
\*\*\*\*\*

PCVPK-15  
IMP999-ECO  
IMP1010-ST  
IMP1011-48  
IMP1011-48

ATATCATGGAGAAGTTGTGTTTGGATGATTTTATGGCTGGTTACCTTGGGATGA  
TTACCATGGTGAAGAAGTGTGTTATTGATGACTTTTATGGCTGGCTGCCGTGGGATGA  
TTACCATGGTGAAGAAGTGTGTTATTGATGACTTTTATGGCTGGCTGCCGTGGGATGA  
TTACCATGGTGAAGAAGTGTGTTATTGATGACTTTTATGGCTGGCTGCCCTGGGATGA  
TTACCATGGTGAAGAAGTGTGTTATTGATGACTTTTATGGCTGGCTGCCCTGGGATGA  
\*\*\*\*\*

FIG. 5 (cont.)

PCVPK-15 TCTACTGAGACTGTGTGACCGGTATCCATTGACTGTAGAGACTAAAGGGGTACTGTTC  
IMP999-ECO TCTACTGAGACTGTGTGATCGATATCCATTGACTGTAGAGACTAAAGGTGGAACCTGTACC  
IMP1010-ST TCTACTGAGACTGTGTGATCGATATCCATTGACTGTAGAGACTAAAGGTGGAACCTGTACC  
IMP1011-48 TCTACTGAGACTGTGTGATCGATATCCATTGACTGTAGAGACTAAAGGTGGAACCTGTACC  
IMP1011-48 TCTACTGAGACTGTGTGATCGATATCCATTGACTGTAGAGACTAAAGGTGGAACCTGTACC  
\*\*\*\*\* \*\* \*\*\*\*\* \*\* \*\*\*\*\* \*\* \*\*\*\*\* \*\*

PCVPK-15 TTTTGTGCCCCGAGTATTTTGATTACCAGCAATCAGGCCCCCAGGAATGGTACTCCTC  
IMP999-ECO TTTTGTGCCCCGAGTATTTTGATTACCAGCAATCAGACCCCCGTGGAAATGGTACTCCTC  
IMP1010-ST TTTTGTGCCCCGAGTATTTTGATTACCAGCAATCAGACCCCCGTGGAAATGGTACTCCTC  
IMP1011-48 TTTTGTGCCCCGAGTATTTTGATTACCAGCAATCAGACCCCCGTGGAAATGGTACTCCTC  
IMP1011-48 TTTTGTGCCCCGAGTATTTTGATTACCAGCAATCAGACCCCCGTGGAAATGGTACTCCTC  
\*\*\*\*\* \*\* \*\*\*\*\* \*\* \*\*\*\*\* \*\* \*\*\*\*\* \*\*

PCVPK-15 AACTGCTGTCCCAGCTGTAGAAGCTCTCTATCGGAGGATTACTTTCGAATTTGGAA  
IMP999-ECO AACTGCTGTCCCAGCTGTAGAAGCTCTCTATCGGAGGATTACTTTCGAATTTGGAA  
IMP1010-ST AACTGCTGTCCCAGCTGTAGAAGCTCTCTATCGGAGGATTACTTTCGAATTTGGAA  
IMP1011-48 AACTGCTGTCCCAGCTGTAGAAGCTCTTATCGGAGGATTACTTTCGAATTTGGAA  
IMP1011-48 AACTGCTGTCCCAGCTGTAGAAGCTCTTATCGGAGGATTACTTTCGAATTTGGAA  
\*\*\*\*\* \*\* \*\*\*\*\* \*\* \*\*\*\*\* \*\* \*\*\*\*\* \*\*

FIG.5 (cont.)

24/26

PCVPK-15  
 IMP999-ECO  
 IMP1010-ST  
 IMP1011-48  
 IMP1011-48

GACTGCTGAGAACAAATCCACGAGGTACCCGAGCCGATTTGAAGCAGTGACCCACC  
 GAATGCTACAGAACAAATCCACGAGGAA--GGGGCCAGTTCGTACCCCTTTCCTCCCCC  
 GAATGCTACAGAACAAATCCACGAGGAA--GGGGCCAGTTCGTACCCCTTTCCTCCCCC  
 GAATGCTACAGAACAAATCCACGAGGAA--GGGGCCAGTTCGTACCCCTTTCCTCCCCC  
 GAATGCTACAGAACAAATCCACGAGGAA--GGGGCCAGTTCGTACCCCTTTCCTCCCCC  
 GAATGCTACAGAACAAATCCACGAGGAA--GGGGCCAGTTCGTACCCCTTTCCTCCCCC

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PCVPK-15  
 IMP999-ECO  
 IMP1010-ST  
 IMP1011-48  
 IMP1011-48

CTGTGCCCTTTTCCCATATAAAATAAAATTACTGAGTCTTTTTTTGTATCACATCGTAATG  
 ATGCCCTGAATTTCCCATATGAAATAAAATTACTGAGTCTTTTT--TATCACTTCGTAATG  
 ATGCCCTGAATTTCCCATATGAAATAAAATTACTGAGTCTTTTT--TATCACTTCGTAATG  
 ATGCCCTGAATTTCCCATATGAAATAAAATTACTGAGTCTTTTT--TATCACTTCGTAATG  
 ATGCCCTGAATTTCCCATATGAAATAAAATTACTGAGTCTTTTT--TATCACTTCGTAATG  
 ATGCCCTGAATTTCCCATATGAAATAAAATTACTGAGTCTTTTT--TATCACTTCGTAATG

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PCVPK-15  
 IMP999-ECO  
 IMP1010-ST  
 IMP1011-48  
 IMP1011-48

GTTTTTATT-TTTATTTA--TTTA--GAGGGTCTTTTAGGATAAAATCTCTGAATTG  
 GTTTTTATTATTATTCATTAGGGTTTAAGTGGGGTCTTTAAGATTAAATCTCTGAATTG  
 GTTTTTATTATTATTCATTAGGGTTTAAGTGGGGTCTTTAAGATTAAATCTCTGAATTG  
 GTTTTTATTATTATTCATTAAAGGGTT-AAGTGGGGGTCCTTAAGATTAAATCTCTGAATTG  
 GTTTTTATTATTATTCATTAAAGGGTT-AAGTGGGGGTCCTTTAAGATTAAATCTCTGAATTG

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 \*\*\*\*\*  
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FIG.5 (cont.)



25/26

TACATAAATAGTCAGCCTTACCACATAATTTGGGCTGTGGCTGC-ATTTTGGAGCGCAT  
 TACATACATGGTTACACGGATATTGTAGTCCCTGG-TCGTATATACTGTTTTCGAACGCAG  
 TACATACATGGTTACACGGATATTGTAGTCCCTGG-TCGTATTTACTGTTTTCGAACGCAG  
 TACATACATGGTTACACGGATATTGTATTCCTGG-TCGTATATACTGTTTTCGAACGCAG  
 TACATACATGGTTACACGGATATTGTATTCCTGG-TCGTATATACTGTTTTCGAACGCAG

\*\*\*\*\* \*\* \* \* \* \* \* \* \* \* \* \* \* \* \* \* \* \* \*

AGCCGAGGCCCTGTGTGCTCGACATTGGTGTGGGTATTTAAATGGAGCCACAGCTGGTTTC  
 TGCCGAGGCCCTACGTGGTCCACATTTCTAGAGGTTTGTAGCCTCAGCCAAAGCTGATTC  
 CGCCGAGGCCCTACGTGGTCCACATTTCCAGAGGTTTGTAGTCTCAGCCAAAGCTGATTC  
 TGCCGAGGCCCTACGTGGTCTACATTTCCAGCAGTTTGTAGTCTCAGCCACAGCTGGTTTC  
 TGCCGAGGCCCTACGTGGTCTACATTTCCAGTAGTTTGTAGTCTCAGCCACAGCTGATTC

\*\*\*\*\* \*\* \* \* \* \* \* \* \* \* \* \* \* \* \* \* \* \* \*

TTTTATTATTGGGTGGAACCAATCAATTGTTGGTCCAGCTCAGGTTTGGGGGTGAAGT  
 TTTTGTATTATTGGTTGGAAGTAATCAATAGTGGAGTCAAGAACAGGTTTGGGTGAAGT  
 TTTTGTATTATTGGTTGGAAGTAATCAATAGTGGAGTCAAGAACAGGTTTGGGTGAAGT  
 TTTTGTGTGTGGTTGGAAGTAATCAATAGTGGAACTAGGACAGGTTTGGGGGTAAAGT  
 TTTTGTGTGTGGTTGGAAGTAATCAATAGTGGAACTAGGACAGGTTTGGGGGTAAAGT

\*\*\*\* \*\* \*\*\*\*\* \*\*\*\*\* \*\* \*\* \*\*\*\*\* \*\*\*\*\* \*\* \*\*\*\*\*

FIG.5 (cont.)

PCVPK-15

IMP999-ECO

IMP1010-ST

IMP1011-48

IMP1011-48

PCVPK-15

IMP999-ECO

IMP1010-ST

IMP1011-48

IMP1011-48

PCVPK-15

IMP999-ECO

IMP1010-ST

IMP1011-48

IMP1011-48

Accession	Sequence
PCVPK-15	GGTCATAGGCCAAGTTGGTGGAGGGGTTACAAAGTTGGCATCCAAGATAACAACAGTGG
IMP999-ECO	GGTCATAGGTTAGGGCTGTGGCCTTTTGTTACAAAGTTATCATCTAGAAATAACAGCAGTGG
IMP1010-ST	GGTCATAGGTTAGGGCTGTGGCCTTTTGTTACAAAGTTATCATCTAGAAATAACAGCAGTGG
IMP1011-48	GGTCATAGGTGAGGGCTGTGGCCTTTTGTTACAAAGTTATCATCTAGAAATAACAGCAGTGG
IMP1011-48	GGTCATAGGTGAGGGCTGTGGCCTTTTGTTACAAAGTTATCATCTAGAAATAACAGCAGTGG

PCVPK-15	ACCCAACACCTCTTTGATTAGAGGTGATGGGGTCTCTGGGGTAA
IIMP999-ECO	AGCCCACTCCCCCTATCACCCCTGGGTGATGGGGGAGCAGGGCCCAG
IIMP1010-ST	AGCCCACTCCCCCTATCACCCCTGGGTGATGGGGGAGCAGGGCCCAG
IIMP1011-48	AGCCCACTCCCCCTGTACACCCTGGGTGATCGGGGAGCAGGGCCCAG
IIMP1011-48	AGCCCACTCCCCCTGTACACCCTGGGTGATCGGGGAGCAGGGCCCAG

FIG. 5 (cont. and end)

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## INTERNATIONAL APPLICATION PUBLISHED UNDER THE PATENT COOPERATION TREATY (PCT)

<b>(51) International Patent Classification <sup>7</sup> :</b> <b>A61K 39/295</b>	<b>A3</b>	<b>(11) International Publication Number:</b> <b>WO 00/01409</b> <b>(43) International Publication Date:</b> 13 January 2000 (13.01.00)
<b>(21) International Application Number:</b> PCT/EP99/04698 <b>(22) International Filing Date:</b> 28 June 1999 (28.06.99) <b>(30) Priority Data:</b> 98/08777                      6 July 1998 (06.07.98)                      FR <b>(71) Applicants (for all designated States except US):</b> MERIAL [FR/FR]; 17, rue Bourgelat, F-69002 Lyon (FR). THE QUEEN'S UNIVERSITY OF BELFAST [GB/GB]; Stoney Road, Stormont, Belfast BT4 3SD (GB). UNIVERSITY OF SASKATCHEWAN [CA/CA]; 52 Campus Drive, Saskatoon, Saskatchewan S7W 5B4 (CA). <b>(72) Inventors; and</b> <b>(75) Inventors/Applicants (for US only):</b> ALLAN, Gordon, Moore [GB/GB]; 51 Cabinhill Gds, Belfast BT5 7AR (GB). MEEHAN, Brian, Martin [GB/GB]; 26 St John's Close, 2 Laganbank Road, Belfast BT1 3LX (GB). ELLIS, John, Albert [US/CA]; 812, 13th Street East, Saskatoon, Saskatchewan S7N 0M3 (CA). KRAKOWKA, George, Steven [US/US]; 2676 Summit Street, Columbus, OH 43202 (US). AUDONNET, Jean-Christophe, Francis [FR/FR]; 119, rue de Créqui, F-69006 Lyon (FR).		<b>(74) Agent:</b> MONCHENY, Michel; Cabinet Lavoix, 9, place d'Estienne d'Orves, F-75441 Paris Cedex 09 (FR). <b>(81) Designated States:</b> AE, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, CA, CH, CN, CU, CZ, DE, DK, EE, ES, FI, GB, GD, GE, GH, GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MD, MG, MK, MN, MW, MX, NO, NZ, PL, PT, RO, RU, SD, SE, SG, SI, SK, SL, TJ, TM, TR, TT, UA, UG, US, UZ, VN, YU, ZA, ZW, ARIPO patent (GH, GM, KE, LS, MW, SD, SL, SZ, UG, ZW), Eurasian patent (AM, AZ, BY, KG, KZ, MD, RU, TJ, TM), European patent (AT, BE, CH, CY, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE), OAPI patent (BF, BJ, CF, CG, CI, CM, GA, GN, GW, ML, MR, NE, SN, TD, TG).  <b>Published</b> <i>With international search report.</i> <b>(88) Date of publication of the international search report:</b> 29 June 2000 (29.06.00)
<b>(54) Title:</b> PORCINE CIRCOVIRUS AND PARVOVIRUS VACCINE		
<b>(57) Abstract</b> <p>The invention relates to antigenic preparations and vaccines directed against the porcine multisystemic wasting syndrome (PMWS), comprising at least one porcine circovirus antigen, preferably type II, and at least one porcine parvovirus antigen.</p>		

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PCT/EP 99/04698

According to International Patent Classification (IPC) or to both national classification and IPC

**Minimum documentation searched (classification system followed by classification symbols)**

IPC 7 A61K

Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched

Electronic data base consulted during the international search (name of data base and, where practical, search terms used)

### C. DOCUMENTS CONSIDERED TO BE RELEVANT

Category *	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
A	<p>DATABASE WPI  Section Ch, Week 9529  Derwent Publications Ltd., London, GB;  Class B04, AN 95-222945  XP002099703  &amp; SU 1 538 305 A (VETERINARY PREPARATIONS  RES INST), 15 December 1994 (1994-12-15)  abstract</p>	1,4,8,9, 16
A	<p>WO 98 03658 A (BAUDU PHILIPPE ;MERIAL  (FR); RIVIERE MICHEL (FR); AUDONNET JEAN  CH) 29 January 1998 (1998-01-29)  cited in the application  page 2, line 16 - line 31; claims 1,13  page 4, line 12 -page 5, line 18</p> <p style="text-align: center;">-/-</p>	1,4,5, 8-10,16

☒ Further documents are listed in the continuation of box C.

☒ Patent family members are listed in annex.

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Date of the actual completion of the international search

31 March 2000

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Authorized officer

Charles, D

## INTERNATIONAL SEARCH REPORT

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PCT/EP 99/04698

## C. (Continuation) DOCUMENTS CONSIDERED TO BE RELEVANT

Category *	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
A	B.M. MEEHAN ET AL.: "CHARACTERIZATION OF NOVEL CIRCOVIRUS DNAs ASSOCIATED WITH WASTING SYNDROMES IN PIGS" JOURNAL OF GENERAL VIROLOGY, vol. 79, no. 9, 1998, pages 2171-2179, XP002099702 the whole document	1,3,4

# INTERNATIONAL SEARCH REPORT

Information on patent family members

International Application No

PCT/EP 99/04698

Patent document cited in search report		Publication date	Patent family member(s)	Publication date
SU 1538305	A	15-12-1994	NONE	
WO 9803658	A	29-01-1998	FR 2751224 A	23-01-1998
			AU 3699197 A	10-02-1998
			CN 1225684 A	11-08-1999
			EP 0912743 A	06-05-1999
			PL 331249 A	05-07-1999

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